CGCGTTTGGTTGCTCGCTCCACCCCGGAGACCTGGTGTGGTGGAGAAATTTGAA CCCGCAGCCTTAGCTCCGAAAAGGCCGAGTTACCTGGCTCTCCCTGAGTGTCGAG GAGGACATGAGTGAAATGACCAGCGAACTCATTTTTATAGGACTCGGTGAAGC CGGATTCTGCATTTCCCTACTTGTAGACTCATTTTGTGGAATAGAGTTGATCGCTG 5 TCTCCTCCGCAAAGCATTTTAACTCGAATAAGCAAATGCCGCCTCTGTTTGAACG TTTTGGTATTTACAAGAGAAATCATTTTACCTAAGAGAACTAATTGAATTGGC CTGCGAAATAGCAGACGGAGAAATTCCTTTGGAAGTTATTCCGTAGCATAAGAG CTGAAACTTCAGAGCAAGTTTTCATTGGGCAAAATGGGGGAACAACCTATCTTCA 10 GCACTCGAGCTCATGTCTTCCAAATTGACCCAAACACAAAGAAGAACTGGGTAC CCACCAGCAAGCATGCAGTTACTGTGTCTTATTTCTATGACAGCACAAGAAATGT CCCAAACATGACATTTACTAAAACATCTCAGAAGTTTGGCCAGTGGGCTGATAGC CGGGCAAACACCGTTTATGGATTGGGATTCTCCTCTGAGCATCATCTTTCGAAAT 15 TTGCAGAAAAGTTTCAGGAATTTAAAGAAGCTGCTCGACTAGCAAAGGAAAAAT CACAAGAGAAGATGGAACTTACCAGTACACCTTCACAGGAATCCGCAGGCGGG ATCTTCAGTCTCCTTTAACACCGGAAAGTATCAACGGGACAGATGATGAAAGAA CACCTGATGTGACACAGAACTCAGAGCCAAGGGCTGAACCAACTCAGAATGCAT TGCCATTTTCACATAGTTCAGCAATCAGCAAACATTGGGAGGCTGAACTGGCTAC 20 CCTCAAAGGAAATAATGCCAAACTCACTGCAGCCCTGCTGGAGTCCACTGCCAAT GTGAAACAATGGAAACAGCAACTTGCTGCCTATCAAGAGGAAGCAGAACGTCTG CACAAGCGGGTGACTGAACTTGAATGTGTTAGTAGCCAAGCAAATGCAGTACAT ACTCATAAGACAGAATTAAATCAGACAATACAAGAACTGGAAGAGACACTGAAA CTGAAGGAAGAGGAATAGAAAGGTTAAAACAAGAAATTGATAATGCCAGAGA 25 ACTACAGAACAGAGGGATTCTTTGACTCAGAAACTACAGGAAGTAGAAATTCG GAACAAGACCTGGAGGGACAACTGTCTGACTTAGAGCAACGTCTGGAGAAAAG TCAGAATGAACAAGAAGCTTTTCGCAATAACCTGAAGACACTCTTAGAAATTCTG GATGGAAAGATATTTGAACTAACAGAATTACGAGATAACTTGGCCAAGCTACTA 30 NNNNNNTTGAATATCACTCCTCCAGGAGGAGGATCTTTTGAAATTGGAATTGTA TATTTCACTGTAAATTTTAGAATCCAGCTTGTAGCTAGTTGGGGAAAAAAAGATGA AAAACTTGAACTACAAATTACCTCCATGTATATTATTGGCCATAGTTAACTAGAA AGTTATAAATAGACACTTAATGCAATCTTTTTTCCTGATATTAGCCAATGGGAGA 35 ATTAACAATGTCTAGGTCACATCCCCTTTTTGTGTTCAACACAGTGAAGATTATCT GCTTTTTAAATTAATTTACGATATCTAGAGCTGTGTTTTTGTGCAAAAACTTA GTGATGAAAGCCTGTCTTTTGTTGTAATCTGAATAATTTCTCAGGATATTTTTGCA TGTATCTTTAATTGAAATATACTATAACTGGGTGTATAGAGTTCTTCCCTTTTTTG 40 TGCTGGAAGATATTTCACTCTGGTGACTACTCTGGTACACTCTGGTGTTCTCTAAT CTTGTCTGTTGTATAGTTTACTTTTCCATATTGATTCCATGTATTTATGAGAAGAT ATTGTCTCCCATTTTATTACACATTTTAAAGCCAACTAACGAAGGCAGCTGAGTC CCTCAGAAATTTTCTTTTAAGTTTCTAATAAATTTGACACACAGTACTGAAATA CAGCAGCCGTCATTGACAGGCTGGTCTAGCAATGTTAAGTATATTTACAGAATA 45 TGCAGTTACATTTATTATATATTTTGCAAGAAATCTTTTCTGAATGATCAATGCA TTTCAATTACGAATAATAATGGTTATTGGGGAACTGTTTATTATAGATAATTTTA AGGTGTATAGCTATTTTAAAGGGGGTCCATTTACATCAAACAGCCGATCAGAGG ACTCTATCTAAATTGTGATCGTGGCAGATAGAGATGGAGTCATGTACTCTATCTG GCTCTACACATCAATCACATCTTGATTCAAACCTCACAAGGCAATATTCTGAATT

GTTAACTAGGTATTTCAAAACAGGAATTAAATTCAATAGGCTCTTCTCAGTGAAC AGGTTTTAATGTTGTTTTGATGTAATTTTAAAAGACTTTTAGCAAACATGCATTTC TTTATATGATATTTCTTTTACGAAGCTATTTTAAAAAGTAAGCCAAGTGCTGTCT AGTCTGCTTATAAAGTAGGAATTGCATCAGAGTACATATATTCTTGCTGTACAAT 5 GCCTGTGATGTTGAGGAGGGTTCTTTTTTAAAGTGTATGCTTGAGTAACTGACTCT ATGGAGTCTATAAATGCACTGACTTCTTGTTTGTACCCCAAAATGATCGAATTGT TAAGTACAAAATTAAGCTAATTAACCAATTTGTAACCATTTTTCACTCATAAAC 10 CTTAAGTGAGTTTTCAGGTGTCTCTGAAAAATTTATAACAATCATGTATTATATGT GCTGTAACATCATGTACGTTACCTCCATCTATTTTAGGATATTTTCCTCACCTATA TATTATAGGGAGAATAATTTAGATACACATGCTCAGAGCTGAGATATTTCTCTGA TAAATCAGGTAACAAAATGTATTTGATTGATGGAATTTTGAAGTAAATGTGTTTT TATCCATCAGTTTCTGAGTAACAAAGAGCACCAAGTTTTAATTTAAATAGGAGAT 15 TTAACACTAGGGATCAGGGAGTTTAGTATGAAGAGTTAAAAAAATTTAAAAAAC AGTGTAAGCTGTTGAAATGGCAAGTGAATTATTTTAATGATGTAATAAAATATTT TTAAATTTTGACATAGTGATCATTTAATGAAAAAACTCACCAAAATGTCTCCATT TGAATTGTATTGATAATGTGGGACATATGTGTGATTCAATATACATATACCCA TATGTATATACAGAAAATTATTTTTAATACTTTCCTACTGATAATGAAATTTAAAA 20 TTGGAAATTTTGTGAGTGTTTTTCTTGTCCAATAGAGCCTAATTGTTTCCTTTTTTA GTGATTTAACAATCTCTTGAGGGCTGCACCTTTAAATTCCCAGATTGTCAATAGA CTTAAGACTTTAACTATTCATTTACAGTAGGAGAGTATGTAGAAATCATCATCC ACAGTCATAATTAGGTTGTGTGCCTACTGTAGTTTTTCCATTTCTGTATTATAT 25 AAACATTTGCATATTAAAATTTGATTTTTCCCAGAGACAAGTATTATATACTGTAT CTATATTTAAATCAAACTGTGGTAATATATTTCTCAGAAAATAATGTTGGGGACT ATAGCCTGAACATGTGGACTTGAAGCGACATGGAGGAGGAGGTTGATCCCATTG TGTATAAGTTAATATGTGATAACTATTGAATCTTGTACAAAAAACAAAAATTGANA AANANAAGAAAAGCAAAAATACAGTTTTTATTTTGAAATACATTTGTTCTCTGG 30 AGAATGTACTTTATCTTTTTCCTCCAGTCTTTTACAGATATTTAAAAGCATTTA AATGATGACAGCATTTACTTAAATCTTTCAGGTGCTACTGGATTTTGCATTAGTGT GTTATGTTGTGAAATCCTAACTTTGACATAAAAGGTTTTATAAGTATTCCCCTGCC TGGAAAATTAGTTTTTCTCCNCTCTCTCTCTTTTCTCTTTTTCTG CAGACTAAAACATGCTCACGAAGTTGCATCTCTCTCTTGTCTCTATAGAAGATCTC 35 CAGCACCATCATAGATTTGATGTTCTGCTGTCATTGNACTGTTGGGAAGCAGTTA GAGGAAAAGCTCACTTTTTTTTCAGGTGGAAATAAAAGGAACACTCAAAATTA AGCCAACACCACCACTACCTTTAAAAACTAGTTTATTTGCCCTGTTAAAATTAAA TGATTCTTNAACATGTGGGCTACAGTCTCCCATGTTTTTATTTAACTGAAGCATAT ACACTTCGGNCATTTATCTCCTGTGGNCCTGATTTTGTCAGTACTGGAATG

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SEQ ID NO: 654
>21590 BLOOD INCYTE_3985758H1
GCNACGGTTGGCGCTCGNCCTGGAGCCTGCCCTGGCGTNCCCCCGCGGGCGCAG
CCAAGCTTCTTGGCNATGGTAGATAACTGCAGGGGACTCTGGCCGCGGCTAACTA
NCCTGGAGATGCTGATCGGGACCCCCCCGCAGAAGCTACAGATTCTCGTTGACA
NTGGAAGCAGTAACTTTGA

SEQ ID NO: 655

>21591 BLOOD 404604.3 AF122922 g4585369 Human Wnt inhibitory factor-1 mRNA, complete cds. 0

- 15 CTGAAGGCAACACCATTCTCCAAACACCTCAAAATGCTATCTTCTTTAAAACATG
 TCAACAAGCTGAGTGCCCAGGCGGGTGCCGAAATGGAGGCTTTTGTAATGAAAG
 ACGCATCTGCGAGTGTCCTGATGGGTTCCACGGACCTCACTGTGAGAAAGCCCTT
 TGTACCCCACGATGTATGAATGGTGGACTTTGTGTGACTCCTGGTTTCTGCATCTG
 CCCACCTGGATTCTATGGAGTGAACTGTGACAAAGCAAACTGCTCAACCACCTGC
 20 TTTAATGGAGGGACCTGTTTCTACCCTGGAAAATGTATTTGCCCTCCAGGACTAG
- 20 TTTAATGGAGGACCTGTTTCTACCCTGGAAAATGTATTTGCCCTCCAGGACTAG AGGGAGAGCAGTGTGAAATCAGCAAATGCCCACAACCCTGTCGAAATGGAGGTA *AATGCATTGGTAAAAGCAAATGTAAGTGTTCCAAAGGTTACCAGGGAGACCTCT !GTTCAAAGCCTGTCTGCGAGCCTGGCTGGCACATGGAACCTGCAATAAAAGGTA
- TTTAAGTTTTCTAAGTACGTCTGTAGCATGATGGTATAGATTTTCTTGTTTCAGTG CTTTGGGACAGATTTTATATTATGTCAATTGATCAGGTTAAAAATTTTCAGTGTGTA GTTGGCAGATATTTTCAAAATTACAATGCATTTATGGTGTCTTGGGGGCAGGGGAA CATCAGAAAGGTTAAATTGGGCAAAAATGCGTAAGTCACAAGAATTTGGATGGT GCAGTTAATGTTGAAGTTACAGCATTTCAGATTTTATTGTCAGATATTTAGATGTT
- 40 GCTTTAGTTTCTGAGCATTGTGTGGAGGTNANCTTTGCACATGCTATCTTATGAA AATAAAATTGGTTGCAATTTAGTGGT

SEQ ID NO: 656

- >21600 BLOOD 480735.6 U60477 g1575342 Human apolipoprotein AI regulatory protein-1/chicken ovalbumin upstream promoter transcription factor II (TFCOUP2) gene, complete cds. 0
 - CATCGAGTGCGTGTGCGGAGACAAGTCGAGCGCAAGCACTACGGCCAGTT CACGTGCGAGGGCTGCAAGAGCTTCTTCAAGCGCAGCGTGCGGAGGAACCTGAG CTACACGTGCCGCCCAACCGGAACTGTCCCATCGACCAGCACCATCGCAACCA

SEQ ID NO: 658 >21621 BLOOD 253228.8 Incyte Unique

- 25 CAGAGCCTGGCCTGGGAGCCAGGATGGCCATCCACAAAGCCTTGGTGATGTGCC
 TGGGACTGCCTCTCTTCCTGTTCCCAGGGGCCTGGGCCCAGGGCCATGTCCCACC
 CGGCTGCAGCCAAGGCCTCAACCCCCTGTACTACAACCTGTGTGACCGCTCTGGG
 GCGTGGGGCATCGTCCTGGAGGCCGTGGCTGGGGGCATTGTCACCACGTTTG
 TGCTCACCATCATCCTGGTGGCCAGCCTCCCCTTTGTGCAGGACACCAAGAAACG
 GAGCCTGCTGGGGACCCAGGTATTCTTCCTTCTGGGGACCCTGGGCCTCTTCTGC
- GAGCCTGCTGGGGACCCAGGTATTCTTCCTTCTGGGGACCCTGGGCCTCTTCTGC
 CTCGTGTTTGCCTGTGGTGAAGCCCGACTTCTCCACCTGTGCCTCTCGGCGCTT
 CCTCTTTGGGGTTCTGTTCGCCATCTGCTTCTCTTGTCTGGCGGCTCACGTCTTTGC
 CCTCAACTTCCTGGCCCGGAAGAACCACGGGCCCCGGGGCTGGTGATCTTCACT
 GTGGCTCTGCTGCTGACCCTGGTAGAGGTCATCAATACAGAGTGGCTGATCA
- 40 GCAGCACAACAGTCCCACCTGGGATGACCCCACGCTGGCCATCGCCCTCGCCGCC
 AATGCCTGGGCCTTCGTCCTCTTCTACGTCATCCCCGAGGTCTCCCAGGTGACCA
 AGTCCAGCCCAGAGCAAAGCTACCAGGGGGACATGTACCCCACCCGGGGCGTGG
 GCTATGAGACCATCCTGAAAGAGCAGAAGGGTCAGAGCATGTCGTGGAGAACA
 AGGCCTTTTCCATGGATGAGCCGGTTGCAGCTAAGAGGCCGGTGTCACCATACAG
- 45 CGGGTACAATGGGCAGCTGCTGACCAGTGTGTACCAGCCCACTGAGATGGCCCT GATGCACAAAGTTCCGTCCGAAGGAGCTTACGACATCATCCTCCCACGGGCCACC GCCAACAGCCAGGTGATGGGCAGTGCCAACTCGACCCTGCGGGCTGAAGACATG TACTCGGCCCAGAGCCACCAGGCGGCCACACCGCCGAAAGACGGCAAGAACTCT CAGGTCTTTAGAAACCCCTACGTGTGGGACTGAGTCAGCGGTGGCGAGGAGAGG

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10 **SEQ ID NO: 659** >21628 BLOOD 255990.10 AJ011497 g4128014 Human mRNA for Claudin-7. 0 GCCGGAGGGGACAGTGGTAGGTGGGGAGGTTGAGTGCAAAGGGTTCAGGCTGTA AGTCATGTTGGGTTGGAATGGGGGCACAGGAAGGTGGGGCTGTTGGGGAGCCAC GCTAAGCCGGGTGTCTGTAGCAGAGCCAGAGAACCGGGACACTGAAGAGGGTGC 15 TGAAGGGGGCGACTCTCAGGGATCGAGCCAGGGCCCCCGAAGGTGGGATCGACC AGGGTAGGAGACAGGAAAAAAAAGGAGAGCAGCGGGTGGGGGCGAAAGCAGG GCCGAGGAGAGACACTTTGGACAGAACCCGGCGGGAAAGGGCGCGCCGAG GCTTGTCAGGGGCGCCCCGCAGCGTCCCAGGCGCACCTGTTGGGAAGAAAGGAA GGGGCTTCCCGGTGTTCGAGGGAAATCCAGTCCGGAGGGGCTGACTCGGAGCTT 20 GGGACTCCTGGGGAGCCACCGCCTCCTCCCAGCGGCGGTCAAAACCGGGCAAG CGAAGGGGCGTGACCCTGGTGCTCAGGTTTCTTCCTCCTCACCTGGGCAAGGAGG GGTGGGGCCACGACTTCCGGTTCAGGTGAGTGTCCCTTCGGTGACGTCAGGTCA ATCCTCGGCCGCCCTCCGGTCCCGCCTCCCCTCCCGCGCTCCCGGGGCGCGCGG 25 CCTGCTGGCTCACCTCCGAGCCACCTCTGCTGCGCACCGCAGCCTCGGACCTACA GCCCAGGATACTTTGGGACTTGCCGGCGCTCAGAAACGCGCCCAGACGGCCCCT CCACCTTTTGTTTGCCTAGGGTCGCCGAGAGCGCCCGGAGGGAACCGCCTGGCCT TCGGGGACCACCATTTTGTCTGGAACCACCCTCCGGCGTATCCTACTCCCTGT GCCGCGAGGCCATCGCTTCACTGGAGGGGTCGATTTGTGTGTAGTTTGGTGACAA GATTTGCATTCACCTGGCCCAAACCCTTTTTGTCTCTTTTGGGTGACCGGAAAACTC 30 GGTCTCCCGCCGGCGCCCCCAGTGTTTTCTGAGGGCGGAAATGGCCAATTCGG CCTGCACCGCCATCCCGCAGTGGCAGATGAGCTCCTATGCGGGTGACAACATCAT 35 CACGCCCAGGCCATGTACAAGGGGCTGTGGATGGACTGCGTCACGCAGAGCAC GGGGATGATGAGCTGCAAAATGTACGACTCGGTGCTCGCCCTGTCCGCGGCCTTG CAGGCCACTCGAGCCCTAATGGTGGTCTCCCTGGTGCTGGGCTTCCTGGCCATGT TTGTGGCCACGATGGGCATGAAGTGCACGCGCTGTGGGGGAGACGACAAAGTGA AGAAGGCCCGTATAGCCATGGGTGGAGGCATAATTTTCATCGTGGCAGGTCTTGC 40 CGCCTTGGTAGCTTCCTGGTATGGCCATCAGATTGTCACAGACTTTTATAACC CTTTGATCCCTACCAACATTAAGTATGAGTTTGGCCCTGCCATCTTTATTGGCTGG GCAGGGTCTGCCCTAGTCATCCTGGGAGGTGCACTGCTCTCTGTTCCTGTCCTG GGAATGAGAGCAAGGCTGGGTACCGTGTACCCCGCTCTTACCCTAAGTCCAACTC TTCCAAGGAGTATGTGTGACCTGGGATCTCCTTGCCCCAGCCTGACAGGCTATGG 45 GAGTGTCTAGATGCCTGAAAGGGCCTGGGGCTGAGCTCAGCCTGTGGGCAGGGT GCCGGACAAAGGCCTCCTGTCACTCTGTCCCTGCACTCCATGTATAGTCCTCTT GGGTTGGGGGGGGGGTGCCGTTGGTGGGAGAGACAAAAAGAGGGAGAGTG TGCTTTTTGTACAGTAATAAAAAATAAGTATTGGGAAGCAGGCTTTTTTCCCTTC AGGGCCTCTGCTTCCTCCCGTCCAGATCCTTGCAGGGAGCTTGGAACCTTAGTG

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SEQ ID NO: 660 >21631 BLOOD 370788.1 AK000072 g7019922 Human cDNA FLJ20065 fis, clone COL01613, highly similar to ECLC_BOVIN EPITHELIAL CHLORIDE CHANNEL PROTEIN. 0

- CTGACCTTCTACTTGGAAAAAAAAAAAATGAATATGGACCACCAGGCAAACTG
 TTTGTCCATGAGTGGGCTCACCTCCGGTGGGGAGTGTTTGATGAGTACAATGAAG

 20 ATCAGCCTTTCTACCGTGCTAAGTCAAAAAAAAATCGAAGCAACAAGGTGTTCCGC
 AGGTATCTCTGGTAGAAAATAGAGTTTATAAGTGTCAAGGAGGAGCAGCTGTCTTAGT
- - 25 CAAAACATAAAGTGCAATTTTAGAAGTACATGGGAGGTGATTAGCAATTCTGAG GATTTTAAAAACACCATACCCATGGTGACACCACCTCCTCCACCTGTCTTCTCATT GCTGAAGATCAGTCAAAGAATTGTGTGCTTAGTTCTTGATAAGTCTGGAAGCATG GGGGGTAAGGACCGCCTAAATCGAATGAATCAAGCAGCAAAACATTTCCTGCTG CAGACTGTTGAAAAATGGATCCTGGGTGGGGATGGTTCACTTTGATAGTACTGCCA

 - 35 AGTAATAGAGATGAGCAAGATAACAGGAGGAAGTCATTTTTATGTTTCAGATGA AGCTCAGAACAATGGCCTCATTGATGCTTTTGGGGCTCTTACATCAGGAAATACT GATCTCTCCCAGAAGTCCCTTCAGCTCGAAAGTAAGGGATTAACACTGAATAGTA ATGCCTGGATGAACGACACTGTCATAATTGATAGTACAGTGGGAAAGGACACGT TCTTTCTCATCACATGGAACAGTCTGCCTCCCAGTATTTCTCTCTGGGATCCCAGT
 - 40 GGAACAATAATGGAAAATTTCACAGTGGATGCAACTTCCAAAATGGCCTATCTC AGTATTCCAGGAACTGCAAAGGTGGGCACTTGGGCATACAATCTTCAAGCCAAA GCGAACCCAGAAACATTAACTATTACAGTAACTTCTCGAGCAGCAAATTCTTCTG TGCCTCCAATCACAGTGAATGCTAAAAATGAATAAGGACGTAAACAGTTTCCCCA GCCCAATGATTGTTTACGCAGAAAATTCTACAAGGATATGTACCTGTTCTTGGAGC
 - 45 CAATGTGACTGCTTTCATTGAATCACAGAATGGACATACAGAAGTTTTGGAACTT
 TTGGATAATGGTGCAGGCGCTGATTCTTTCAAGAATGATGGAGTCTACTCCAGGT
 ATTTTACAGCATATACAGAAAATGGCAGATATAGCTTAAAAGTTCGGGCTCATGG
 AGGAGCAAACACTGCCAGGCTAAAATTACGGCCTCCACTGAATAGAGCCGCGTA
 CATACCAGGCTGGTAGTGAACGGGGAAATTGAAGCAAACCCGCCAAGACCTGA

AATTGATGAGGATACTCAGACCACCTTGGAGGATTTCAGCCGAACAGCATCCGG AGGTGCATTTGTGGTATCACAAGTCCCAAGCCTTCCCTTGCCTGACCAATACCCA CCAAGTCAAATCACAGACCTTGATGCCACAGTTCATGAGGATAAGATTATTCTTA CATGGACAGCACCAGGAGATAATTTTGATGTTGGAAAAGTTCAACGTTATATCAT 5 AAGAATAAGTGCAAGTATTCTTGATCTAAGAGACAGTTTTGATGATGCTCTTCAA GTAAATACTACTGATCTGTCACCAAAGGAGGCCAACTCCAAGGAAAGCTTTGCA AAAGTATAGATAAAAGCAATTTGACATCAAAAGTATCCAACATTGCACAAGTAA CTTTGTTTATCCCTCAAGCAAATCCTGATGACATTGATCCTACACCTACTTCCTAC 10 TCCTACTCCTACTCCTGATAAAAGTCATAATTCTGGAGTTAATATTTCTACGCTGG TATTGTCTGTGATTGGGTCTGTTGTAATTGTTAACTTTATTTTAAGTACCACCATT TGAACCTTAACGAAGAAAAAATCTTCAAGTAGACCTAGAAGAGAGTTTTAAAA AACAAAACAATGTAAGTAAAGGATATTTCTGAATCTTAAAATTCATCCCATGTGT GATCATAAACTCATAAAAATAATTTTAAGATGTCGGAAAAGGATACTTTGATTAA 15 ATAAAAACACTCATGGATATGTAAAAACTGTCAAGATTAAAATTTAATAGTTTCA TTTATTTGTTATTTGTAAGAAATAGTGATGAACAAGATCCTTTTTCATAC TGATACCTGGTTGTATATTTGATGCAACAGTTTTCTGAAATGATATTTCAAAT TGCATCAAGAAATTAAAATCATCTATCTGAGTAGTCAAAATACAAGTAAAGGAG AGCAAATAAACAACATTTGGAAAAAAATG

20

SEO ID NO: 661

25 AAAGCTTATGGCTCTGTGATGATATTAGTGACCAGCGGAGATGATAAGCTTCTTG GCAATTGCTTACCCACTGTGCTCAGCAGTGGTTCAACAATTCACTCCATTGCCCT GGGTTCATCTGCAGCCCCAAATCTGGA

SEQ ID NO: 662

30 >21660 BLOOD 238908.1 AL137516 g6808175 Human mRNA; cDNA DKFZp564M2178 (from clone DKFZp564M2178); partial cds. 0 GAACCACCGGCAGACGCACCTCCGGGCCACACCCACGAGGCTCCTGCCCCTGTT GTCCTGGGGTCCCCAGTTGTTCTAGGGCCTCCTGTGGGCCAGGCCCGAGTGGCTG TGGAGCACTCATACCGAAAGGCAGAAGAGGGTGGGGAAGGGGCGACTGTCCCAT CTGCCGCTGCCACCACCACTGAGGTAGTGACTGAGGTGGAGCTGCTCCTCTACAA 35 GTGCTCTGAGTGCTCCCAGCTCTTCCAGCTGCCGGCGGATTTCCTGGAGCACCAG AGGTGCAGGCCTCGTCACCTGCAGAGGTGCCTGTGTCTCAGCCTGACCCCTTGCC AGCTTCTGACCACAGTTACGAGCTGCGCAATGGTGAAGCCATTGGGCGGGATCG 40 CCGGGGGCCAGGGCCCGGAGGAACACAGTGGAGAAGCAGGCGGGGCAGCCA CACAGGAGCTCTTCTGCTCAGCCTGTGACCAGCTCTTTCTCTCACCCCACCAGCTA CAGCAGCACCTGCGGAGTCACCGGGAGGGCGTCTTTAAGTGCCCCCTGTGCAGTC GTGTCTTCCCTAGCCCTTCCAGTCTGGACCAGCACCTTGGAGACCATAGCAGCGA GTCACACTTCCTGTGTGTAGACTGTGGCCTGGCCTTCGGCACAGAGGCCCTCCTC 45 CTGGCCCACCGGCGAGCCCACACCCCGAATCCTCTGCATTCATGTCCATGTGGGA AGACCTTTGTCAACCTTACCAAGTTCCTTTATCACCGGCGTACTCATGGGGTAGG GGGTGTCCCTCTGCCCACAACACCAGTCCCACCAGAGGAACCTGTCATTGGTTTC CCTGAGCCAGCCCAGCAGAGACTGGAGAGCCAGAGGCCCCTGTG

TCTGAGGAGACCTCAGCAGGGCCCGCTGCCCCAGGCACCTACCGCTGCCTCTGT

GCAGCCGTGAATTTGGAAAGGCCTTGCAGCTGACCCGGCACCAACGTTTTGTGCA TCGGCTGGAGCGCCCATAAATGCAGCATTTGTGGCAAGATGTTCAAGAAGAA GTCTCACGTGCGTAACCACCTGCGCACACACACGGGGAGCGGCCCTTCCCCTGC CCTGACTGCTCCAAGCCCTTCAACTCACCTGCCAACCTGGCCCGCCACCGGCTCA 5 CACACAGGAGAGCGCCCTACCGGTGTGGGGACTGTGGCAAGGCTTTCACGC AAAGCTCCACACTGAGGCAGCACCGCTTGGTGCATGCCCAGCACTTCCCCTACCG $\tt CTGCCAGGAATGTGGGGTGCGTTTTCACCGTCCTTACCGCCTGCTCATGCACCGC$ TACCATCACACAGGTGAATACCCCTACAAGTGTCGCGAGTGCCCCCGCTCCTTCT TGCTGCGTCGGCTGCAGGTGCACCAGCTCGTGGTCCATGCCGGGCGCCAGCC 10 CCACCGCTGCCCATCCTGTGGGGCTGCCTTCCCCTCCTCACTGCGGCTCCGGGAG CACCGCTGTGCAGCCGCTGCTGCCCAGGCCCCACGGCGCTTTGAGTGTGGCACCT GTGGCAAGAAAGTGGGCTCAGCTGCTCGACTGCAGGCACACGAGGCGCCCATG CAGCTGCTGGGCCTGGAGAGGTCCTGGCTAAGGAGCCCCCTGCCCCTCGAGCCCC ACGGGCCACTCGTGCACCAGTTGCCTCTCCAGCAGCCCTTGGAAGCACTGCTACA 15 GCATCCCCTGCGGCCCCGCCGCCGGGGTCTAGAGTGCAGCGAGTGCAAG AAGCTGTTCAGCACAGAGACGTCACTGCAGGTGCACCGGCGCATCCACACAGGT GAGCGCCATACCCATGTCCAGACTGTGGCAAAGCGTTCCGTCAGAGTACCCAC CTGAAAGACCACCGGCGCCTGCACACAGGTGAGCGGCCCTTTGCCTGTGAAGTG TGTGGCAAGGCCTTTGCCATCTCCATGCGCCTGGCAGAACATCGCCGCATCCACA 20 CAGGCGAACGACCCTACTCCTGCCCTGACTGTGGCAAGAGCTACCGCTCCTTCTC CAACCTCTGGAAGCACCGCAAGACCCATCAGCAGCAGCATCAGGCAGCTGTGCG .GCAGCAGCTGGCAGAGGCGGAGGCTGCCGTTGGCCTGGCCGTCATGGAGACTGC TGTGGAGGCGCTACCCCTGGTGGAAGCCATTGAGATCTACCCTCTGGCCGAGGCT · GAGGGGTCCAGATCAGTGGCTGACTCTGCCCGACTTCCTCTTTGGCACCTCCAT 25 TCCCTGTTGCTGAAGGCCCTCCAGCATCCCCTTAAGCATCTGTACATACTGTGTCC CTTCCTCTTCCCATCCCACCACCTTGTAAGTTCTAAATTGGATTTATTCTCTCGT CTCTTAGCACTGGTGACCCCAAAAATGAAACCATCAATAAAGACTGAGTTGCC

30 **SEQ ID NO: 663** >21669 BLOOD 132774.1 Incyte Unique GCCGGACAGAGCAGAAGCCCTCTTGGACTGGACGATTTGGGAATTCAAAACT TGGGACAAACTGTCAGCCTTGCCCCTGCTGTGGAGGCAGCCTCAATGCTGAAAAT GGAGCCTCTGAACAGCACGCACCCGGCACCGCCTCCAGCAGCCCCCTGGA 35 GTCCCGTGCGGCGGTGGCGGCAGCGCAATGGCAACGAGTACTTCTACATTCTG GTTGTCATGTCCTTCTACGGCATTTTCTTGATCGGAATCATGCTGGGCTACATGAA ATCCAAGAGGCGGGAGAAGACTCCAGCCTCCTGCTGCTGTACAAAGACGAGGA GCGGCTCTGGGGGGAGGCCATGAAGCCGCTGCCCGTGGTGTCGGGCCTGAGGTC GGTGCAGGTGCCCCTGATGCTGAACATGCTGCAGGAGAGCGTGGCGCCCGCGCT 40 GTCCTGCACCCTCTGTTCCATGGAAGGGGACAGCGTGAGCTCCGAGTCCTCCTCC CCGGACGTGCACCTCACCATTCAGGAGGAGGGGGGGCAGACGAGGAGCTGGAGGA GACCTCGGAGACGCCCTCAACGAGAGCAGCGAAGGGTCCTCGGAGAACATCCA CTTAGAGAGAGAAAGACAGTTTTCAAGTGTCTGGTTTCACTTTCACAGTGCGGC 45 AGGCTCAGCCGGAACCAGCACCTCCAAGGAGTCCGGGAGGTGCCTGTGGTTTAC ACCCACCACTGAAAAAGCCGCGGAGATGCGCAGCGCGTACACTGACTTTGGGGC CTGGGTGTTGGGGTTCTGATCAGAATTTGGCGGGATGATATGCTTGCCATTTTCTC ACTGGATGCCCTGGGTAGCTCCTGCAGGGTCTGCCTGTTCCCAGGGCTGCCGAAT

GCTTAGGACACGCTGAGAGACTAGTTGTGATTTGCTATTTTGCCTAGAGCTTTGT CCTTCTAGATCTGATTGGCTGTAAGTATCTCTACTGTGTACCTGTGGCATTCCTTC ACAGTGGGTTACAAGCTTCTTTGGGATTAGAGGGGGATTTTGGATGGGAGAAAG CGTGGGAGATCGTGGAACCCCAGCCCCATTTGCACACTATAAGAAAAAAAGTAA CTTTTAAACCTGTTAACATTGGCCGGGGTTATAAGAGATGATCTTCTATTT

SEQ ID NO: 664

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>21683 BLOOD 444662.14 Z58148 g1029379 Human CpG island DNA genomic Mse1 fragment, clone 30a7, forward read cpg30a7.ft1d. 3e-15

- 10 CTCAGCGTCAGGCAAGTTGGCCTCTCTGTTGTAAATTAGTGGTTAAGGTTATCTA CGGGTGATTGGCTTTTTTTTTTTGGCAAACCAGTTATTCAAGTTTCTGGTCTTTAA AAAACTCTGTGGCGGTACGGTAACCGAGGAGGTTCCAGCGCGGCGGAAGTACCC CGCGGGTGGGTGTGCGCAAGGCCAGGGCCAGAGGGCACGTGGCGCCGGGA 15 GGAGAGAGATGTCTTTTCGAGGCGGAGGTCGTGGAGGCTTTAATCGAGGTGGT GGAGGTGGCCGGCTTCAACCGAGGCGGCAGCAACCACTTCCGAGGTGGAGG CGGCGGTGGAGGCGCGGCAATTTCAGAGGCGGCGGCAGGGAGGATTTGGAC GAGGGGTGGCCGCGGAGGCTTTAACAAAGGCCAAGACCAAGGACCTCCAGAA CGTGTAGTCTTATTAGGAGAGTTCCTGCATCCCTGTGAAGATGACATAGTTTGTA 20 AATGTACCACAGATGAAAATAAGGTGCCTTATTTCAATGCTCCTGTTTACTTAGA
- AAACAAAGAACAAATTGGAAAAGTGGATGAAATATTTGGACAACTCAGAGATTT TTATTTTCAGTTAAGTTGTCAGAAAACATGAAGGCTTCATCCTTTAAAAAACTA CAGAAGTTTTATATAGACCCATATAAGCTGCTGCCACTGCAGAGGTTTTTACCTC GACCTCCAGGTGAGAAAGGACCTCCAAGAGGTGGTGGCAGGGGAGGCCGAGGA 25 GGAGGAAGAGGAGGTGGCAGAGGTGGTGGCAGAGGCGGTGGTTTTAGAGG
- TGGAAGAGGAGGTGGAGGTGGGGGCTTCAGAGGAGGAAGAGGTGGTGTTTCA GAGGGAGAGACATTAAGTGAAACAGTTGACAGACATCACCAGTTGACTTCTGC ATTAACCTGCATGATCTGTTTCTACTATGGATTGGAAACTTGTTTCTTGAACAAGT CTTGAAGATCTTGGTCATTTTATGACAATGGATCTAAAATGTCAGCATCATGCAA
- AGTGCAACGGAATAGTGAATTTTGCTCTAAAAGAGCATGAACAAGTCTTTCTAAT 30 GTTTTGTACAGTGCCTGGNACTCTGTGGGTGCTCAATAAATGGATAGGAGTTTTC

SEO ID NO: 665

- 35 yp61a02.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:191882 3', mRNA sequence gi|908298|gb|H38799.1|H38799[908298] TGATGCATCCTAAAATNNTAAGCTTCAAATCTGATTTGGTATCACCGAGGAAACC TTGCCCCCATCACTCAGCATTGCACTTAGATACAGAATGAGTTAGATAAACTTGG CTTGTCTAGAGACCCATGTCATCTTAACCTAAAGGGAAATCTTATTGCGTTATCA 40
- AGTTCTCACTAAAACAATCCTGAGATTTCTTAATTTCATGGGTTCTTTAAATATTA TAAACACAGAGTCAACATAGGAATGAAATTGTATTTGTTAAAATACACACATTG GGGGGNCAAGAGGCAGATGACTACTTTTC

GGAGGTAATGCTTGCTCCTCCNAAAAGGCNGGTTTTCCATCCGGGGG

45

SEQ ID NO: 666

>21694 BLOOD 029567.1 Incyte Unique

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SEQ ID NO: 667 >21697 BLOOD 350207.6 X69086 g34811 Human mRNA for utrophin. 0 GCGGGCAGCAGCCGGCCGCGGGCTTTCTCCCGCCGAGGGGCGAGGAGGAGC ${\tt CTCTGGCTCCAGAAGCCGATTGGGGAATCACGGGGAGCGGCGCCCCCTTCTTTT}$ 20 GGGTCATTTCTGCAAACGGAAAACTCTGTAGCGTTTGGCAAAGTTGGTGCCTGCN CGCCCTTCCAGGTTTGCGCTTTGACTGTTTTGTTTTTGGCGGAACTACCAGGCAG 25 TAAAATACATCGCACCACCAAACTAACACTCGCACACCCCCCGCGGTTACTCCG TGTCAAACTCCTAGAGGAGCCCTTGGCCAGCTCGGGGTGCGGCGGTGGCGACCG GCAGGCGAGGAGCCCGCGGGCAGCAGGTATTGATGTCAAGCTGAACCATCGTA GGAAGTTGAAAGCCTTAGAAAGAGGACTTGGTAAAGTTTTTGGATTATCTTGAAA CTCTGGCAAAATGGCCAAGTATGGAGAACATGAAGCCAGTCCTGACAATGGGCA GAACGAATTCAGTGATATCATTAAGTCCAGATCTGATGAACACAATGACGTACA 30 GAAGAAAACCTTTACCAAATGGATAAATGCTCGATTTTCAAAGAGTGGGAAACC CTAGAAGGCCTCACAGGAACATCACTGCCAAAGGAACGTGGTTCCACAAGGGTA CATGCCTTAAATAACGTCAACAGAGTGCTGCAGGTTTTACATCAGAACAATGTGG AATTAGTGAATATAGGGGGAACTGACATTGTGGATGGAAATCACAAACTGACTT 35 TGGGGTTACTTTGGAGCATCATTTTGCACTGGCAGGTGAAAGATGTCATGAAGGA TGTCATGTCGGACCTGCAGCAGACGAACAGTGAGAAGATCCTGCTCAGCTGGGT GCGTCAGACCACCAGGCCTACAGCCAAGTCAACGTCCTCAACTTCACCACCAGC TGGACAGATGGACTCGCCTTTAATGCTGTCCTCCACCGACATAAACCTGATCTCT 40 TCAGCTGGGATAAAGTTGTCAAAATGTCACCAATTGAGAGACTTGAACATGCCTT CAGCAAGGCTCAAACTTATTTGGGAATTGAAAAGCTGTTAGATCCTGAAGATGTT GGTGCTACCTCAGCAAGTCACCATAGACGCCATCCGTGAGGTAGAGACACTCCC AAGGAAATATAAAAAAGAATGTGAAGAAGAGGCAATTAATATACAGAGTACAG 45 CGCCTGAGGAGGAGCATGAGAGTCCCCGAGCTGAAACTCCCAGCACTGTCACTG AGGTCGACATGGATCTGGACAGCTATCAGATTGCGTTGGAGGAAGTGCTGACCT GGTTGCTTTCTGAGGACACTTTCCAGGAGCAGGATGATATTTCTGATGATGT TGAAGAAGTCAAAGACCAGTTTGCAACCCATGAAGCTTTTATGATGGAACTGACT

CAAGGAACTCTGTCAGACGAAGAAGAATTTGAGATTCAGGAACAGATGACCCTG CTGCACGATGTGCTGATGGAACTGCAGAAGAAGCAACTGCAGCAGCTCTCCGCC TGGTTAACACTCACAGAGGAGCGCATTCAGAAGATGGAAACTTGCCCCCTGGAT 5 GATGATGTAAAATCTCTACAAAAGCTGCTAGAAGAACATAAAAGTTTGCAAAGT GATCTTGAGGCTGAACAGGTGAAAGTAAATTCACTAACTCACATGGTGGTCATTG TTGATGAAAACAGTGGTGAGAGCGCTACAGCTATCCTAGAAGACCAGTTACAGA AACTTGGTGAGCGCTGGACAGCAGTATGCCGTTGGACTGAAGAACGCTGGAATA GGTTACAAGAAATCAATATTGTGGCAGGAATTATTGGAAGAACAGTGCTTGTT 10 CTTCAAAGACCAAAAGGAACTAAGTGTCAGTGTTCGACGTCTGGCTATTTTGAAG GATGTGGGACAATTACTTGATAATTCCAAGGCATCTAAGAAGATCAACAGTGAC TCAGAGGAACTGACTCAAAGATGGGATTCTTTGGTTCAGAGACTAGAAGATTCCT 15 CCAACCAGGTGACTCAGGCTGTAGCAAAGCTGGGGATGTCTCAGATTCCTCAGA AGGACCTTTTGGAGACTGTTCGTGTAAGAGAACAAGCAATTACAAAAAATCTA AGCAGGAACTGCCTCCTCCTCCCCCAAAGAAGAGACAGATCCATGTGGATAT TGAAGCTAAGAAAAGTTTGATGCTATAAGTGCAGAGCTGTTGAACTGGATTTTG AAATGGAAAACTGCCATTCAGACCACAGAGATAAAAGAGTATATGAAGATGCAA 20 GACACTTCCGAAATGAAAAAGAAGTTGAAGGCATTAGAAAAAGAACAGAGAGA AAGAATCCCCAGAGCAGATGAATTAAACCAAACTGGACAAATCCTTGTGGAGCA *AATGGGAAAAGAAGGCCTTCCTACTGAAGAAATAAAAAATGTTCTGGAGAAGGT GTCATCAAGACAAAGGAGGAGTGGGTAAAACACACTTCCATTTCTGAATCTTCCC 25 GGCAGTCCTTGCCAAGCTTGAAGGATTCCTGTCAGCGGGAATTGACAAATCTTCT TGGCCTTCACCCCAAAATTGAAATGGCTCGTGCAAGCTGCTCGGCCCTGATGTCT CAGCCTTCTGCCCCAGATTTTGTCCAGCGGGGCTTCGATAGCTTTCTGGGCCGCT ACCAAGCTGTACAAGAGGCTGTAGAGGATCGTCAACAACATCTAGAGAATGAAC 30 TGAAGGGCCAACCTGGACATGCATATCTGGAAACATTGAAAACACTGAAAGATG TGCTAAATGATTCAGAAAATAAGGCCCAGGTGTCTCTGAATGTCCTTAATGATCT TGCCAAGGTGGAGAAGGCCCTGCAAGAAAAAAAGACCCTTGATGAAATCCTTGA GAATCAGAAACCTGCATTACATAAACTTGCAGAAGAAACAAAGGCTCTGGAGAA AAATGTTCATCCTGATGTAGAAAAATTATATAAGCAAGAATTTGATGATGTGCAA 35 GGAAAGTGGAACAAGCTAAAGGTCTTGGTTTCCAAAGATCTACATTTGCTTGAGG AAATTGCTCTCACACTCAGAGCTTTTGAGGCCGATTCAACAGTCATTGAGAAGTG GATGGATGGCGTGAAAGACTTCTTAATGAAACAGCAGGCTGCCCAAGGAGACGA CGCAGGTCTACAGAGGCAGTTAGACCAGTGCTCTGCATTTGTTAATGAAATAGAA ACAATTGAATCATCTCTGAAAAACATGAAGGAAATAGAGACTAATCTTCGAAGT 40 GGTCCAGTTGCTGGAATAAAAACTTGGGTGCAGACAAGACTAGGTGACTACCAA ACTCAACTGGAGAAACTTAGCAAGGAGATCGCTACTCAAAAAAGTAGGTTGTCT GAAAGTCAAGAAAAGCTGCGAACCTGAAGAAGACTTGGCAGAGATGCAGGA ATGGATGACCCAGGCCGAGGAAGAATATTTGGAGCGGGATTTTGAGTACAAGTC ACCAGAAGAGCTTGAGAGTGCTGTGGAAGAGATGAAGAGGGCAAAAGAGGATG 45 TGTTGCAGAAGGAGGTGAGAGTGAAGATTCTCAAGGACAACATCAAGTTATTAG CTGCCAAGGTGCCCTCTGGTGGCCAGGAGTTGACGTCTGAGCTGAATGTTGTGCT GGAGAATTACCAACTTCTTTGTAATAGAATTCGAGGAAAGTGCCACACGCTAGA GGAGGTCTGGTTGTTGGATTGAACTGCTTCACTATTTGGATCTTGAAACTACCT

CGGATGCTGTCAACGAAGCCCTGGAGTCTCTGGAATCTGTTCTGCGCCACCCGGC CCTGGATGATAATCAGTGAGAAACTGGAGGCTTTCAACAGCCGATATGAAGA TCTAAGTCACCTGGCAGAGAGCAAGCAGATTTCTTTGGAAAAGCAACTCCAGGT 5 GCTGCGGGAAACTGACCAGATGCTTCAAGTCTTGCAAGAGAGCTTGGGGGAGCT GGACAAACAGCTCACCACATACCTGACTGACAGGATAGATGCTTTCCAAGTTCCA CAGGAAGCTCAGAAAATCCAAGCAGAGATCTCAGCCCATGAGCTAACCCTAGAG GAGTTGAGAAGAAATATGCGTTCTCAGCCCCTGACCTCCCCAGAGAGTAGGACT GCCAGAGGAGGAAGTCAGATGGATGTGCTACAGAGGAAACTCCGAGAGGTGTCC 10 ACAAAGTTCCAGCTTTTCCAGAAGCCAGCTAACTTCGAGCAGCGCATGCTGGACT GCAAGCGTGTGCTGGATGGCGTGAAAGCAGAACTTCACGTTCTGGATGTGAAGG ACGTAGACCCTGACGTCATACAGACGCACCTGGACAAGTGTATGAAACTGTATA AAACTTTGAGTGAAGTCAAACTTGAAGTGGAAACTGTGATTAAAACAGGAAGAC 15 CTTCCCTGAAGGTTCTTTACAATGACCTGGGCGCACAGGTGACAGAAGGAAAAC AGGATCTGGAAAGAGCATCACAGTTGGCCCGGAAAATGAAGAAGAGGCTGCTT CTCTCTCTGAATGGCTTTCTGCTACTGAAACTGAATTGGTACAGAAGTCCACTTC AGAAGGTCTGCTTGGTGACTTGGATACAGAAATTTCCTGGGCTAAAAATGTTCTG AAGGATCTGGAAAAGAGAAAAGCTGATTTAAATACCATCACAGAGAGTAGTGCT 20 GCCCTGCAAAACTTGATTGAGGGCAGTGAGCCTATTTTAGAAGAGAGGCTCTGC GTCCTTAACGCTGGGTGGAGCCGAGTTCGTACCTGGACTGAAGATTGGTGCAATA CCTTGATGAACCATCAGAACCAGCTAGAAATATTTGATGGGAACGTGGCTCACAT AAGTACCTGGCTTTATCAAGCTGAAGCTCTATTGGATGAAATTGAAAAGCAAACC AACAAGTAAACAGGAAGAAATTGTGAAGCGTTTAGTATCTGAGCTGGATGATGC : * 25 CAACCTCCAGGTTGAAAATGTCCGCGATCAAGCCCTTATTTTGATGAATGCCCGT GGAAGCTCAAGCAGGGAGCTTGTAGAACCAAAGTTAGCTGAGCTGAATAGGAAC TTTGAAAAGGTGTCTCAACATATCAAAAGTGCCAAATTGCTAATTGCTCAGGAAC CATTATACCAATGTTTGGTCACCACTGAAACATTTGAAACTGGTGTGCCTTTCTCT GACTTGGAAAAATTAGAAAATGACATAGAAAATATGTTAAAATTTGTGGAAAAA 30 CACTTGGAATCCAGTGATGAAGATGAAAAGATGGATGAGGAGAGTGCCCAGATT GAGGAAGTTCTACAAAGAGGAGAAGAAATGTTACATCAACCTATGGAAGATAAT AAAAAAGAAAAGATCCGTTTGCAATTATTACTTTTGCATACTAGATACAACAAA TTAAGGCAATCCCTATTCAACAGAGGAAAATGGGTCAACTTGCTTCTGGAATTAG ATCATCACTTCTTCCTACAGATTATCTGGTTGAAAATTAACAAAATTTTACTTTGCA 35 TGGATGATGTTGAATTATCGCTTAATGTTCCAGAGCTCAACACTGCTATTTACGA AGACTTCTCTTTTCAGGAAGACTCTCTGAAGAATATCAAAGACCAACTGGACAAA CTTGGAGAGCAGATTGCAGTCATTCATGAAAAACAGCCAGATGTCATCCTTGAA GCCTCTGGACCTGAAGCCATTCAGATCAGAGATACACTTACTCAGCTGAATGCAA AATGGGACAGAATTAATAGAATGTACAGTGATCGGAAAGGTTGTTTTGACAGGG 40 CAATGGAAGAATGGAGACAGTTCCATTGTGACCTTAATGACCTCACACAGTGGA TAACAGAGGCTGAAGAATTACTGGTTGATACCTGTGCTCCAGGTGGCAGCCTGG ACTTAGAGAAAGCCAGGATACATCAGCAGGAACTTGAGGTGGGCATCAGCAGCC ACCAGCCAGTTTTGCAGCACTAAACCGAACTGGGGATGGGATTGTGCAGAAAC TCTCCCAGGCAGATGGAAGCTTCTTGAAAGAAAAACTGGCAGGTTTAAACCAAC 45 GCTGGGATGCAATTGTTGCAGAAGTGAAGGATAGGCAGCCAAGGCTAAAAGGAG AAAGTAAGCAGGTGATGAAGTACAGGCATCAGCTAGATGAGATTATCTGTTGGT TAACAAAGGCTGAGCATGCTATGCAAAAGAGATCAACCACCGAATTGGGAGAAA ACCTGCAAGAATTAAGAGACTTAACTCAAGAAATGGAAGTACATGCTGAAAAAC TCAAATGGCTGAATAGAACTGAATTGGAGATGCTTTCAGATAAAAGTCTGAGTTT

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ACCTGAAAGGGATAAAATTTCAGAAAGCTTAAGGACTGTAAATATGACATGGAA TAAGATTTGCAGAGAGGTGCCTACCACCCTGAAGGAATGCATCCAGGAGCCCAG TTCTGTTTCACAGACAAGGATTGCTGCTCATCCTAATGTCCAAAAGGTGGTGCTA GTATCATCTGCGTCAGATATTCCTGTTCAGTCTCATCGTACTTCGGAAATTTCAAT GACCAGATGCTGAAGTCCAACATTGTCACTGTTGGGGATGTAGAAGAGATCAAT AAGACCGTTTCCCGAATGAAAATTACAAAGGCTGACTTAGAACAGCGCCATCCT CAGCTGGATTATGTTTTTACATTGGCACAGAATTTGAAAAATAAAGCTTCCAGTT CAGATATGAGAACAGCAATTACAGAAAAATTGGAAAGGGTCAAGAACCAGTGG GATGGCACCCAGCATGGCGTTGAGCTAAGACAGCAGCAGCTTGAGGACATGATT ATTGACAGTCTTCAGTGGGATGACCATAGGGAGGAGACTGAAGAACTGATGAGA AAATATGAGGCTCGACTCTATATTCTTCAGCAAGCCCGACGGGATCCACTCACCA AACAAATTTCTGATAACCAAATACTGCTTCAAGAACTGGGTCCTGGAGATGGTAT CGTCATGGCGTTCGATAACGTCCTGCAGAAACTCCTGGAGGAATATGGGAGTGA TGACACAAGGAATGTGAAAGAAACCACAGAGTACTTAAAAACATCATGGATCAA TCTCAAACAAGTATTGCTGACAGACAGACGCCTTGGAGGCTGAGTGGAGGAC GGTGCAGGCCTCTCGCAGAGATCTGGAAAACTTCCTGAAGTGGATCCAAGAAGC AGAGACCACAGTGAATGTGCTTGTGGATGCCTCTCATCGGGAGAATGCTCTTCAG GATAGTATCTTGGCCAGGGAACTCAAACAGCAGATGCAGGACATCCAGGCAGAA ATTGATGCCCACAATGACATATTTAAAAGCATTGACGGAAACAGGCAGAAGATG GTAAAAGCTTTGGGAAATTCTGAAGAGGCTACTATGCTTCAACATCGACTGGATG *********ATATGAACCAAAGATGGAATGACTTAAAAGCAAAATCTGCTAGCATCAGGGCCC ATTTGGAGGCCAGCGCTGAGAAGTGGAACAGGTTGCTGATGTCCTTAGAAGAAC 1 1 mais TGATCAAATGGCTGAATATGAAAGATGAAGAGCTTAAGAAACAAATGCCTATTG GAGGAGATGTTCCAGCCTTACAGCTCCAGTATGACCATTGTAAGGCCCTGAGACG GGAGTTAAAGGAGAAAGAATATTCTGTCCTGAATGCTGTCGACCAGGCCCGAGT TTTCTTGGCTGATCAGCCAATTGAGGCCCCTGAAGAGCCAAGAAGAAACCTACA ATCAAAAACAGAATTAACTCCTGAGGAGAGAGCCCAAAAGATTGCCAAAGCCAT GCGCAAACAGTCTTCTGAAGTCAAAGAAAAATGGGAAAGTCTAAATGCTGTAAC TAGCAATTGGCAAAAGCAAGTGGACAAGGCATTGGAGAAACTCAGAGACCTGCA GGGAGCTATGGATGACCTGGACGTGACATGAAGGAGGCAGAGTCCGTGCGGAA TGGCTGGAAGCCCGTGGGAGACTTACTCATTGACTCGCTGCAGGATCACATTGAA AAAATCATGGCATTTAGAGAAGAAATGCACCAATCAACTTTAAAGTTAAAACGG TGAATGATTTATCCAGTCAGCTGTCTCCACTTGACCTGCATCCCTCTCTAAAGATG TCTCGCCAGCTAGATGACCTTAATATGCGATGGAAACTTTTACAGGTTTCTGTGG ATGATCGCCTTAAACAGCTTCAGGAAGCCCACAGAGATTTTGGACCATCCTCTCA GCATTTTCTCTCTACGTCAGTCCAGCTGCCGTGGCAAAGATCCATTTCACATAAT AAAGTGCCCTATTACATCAACCATCAAACACAGACCACCTGTTGGGACCATCCTA AAATGACCGAACTCTTTCAATCCCTTGCTGACCTGAATAATGTACGTTTTTCTGCC TACCGTACAGCAATCAAAATCCGAAGACTACAAAAAGCACTATGTTTGGATCTCT TAGAGTTGAGTACAACAAATGAAATTTTCAAACAGCACAAGTTGAACCAAAATG ACCAGCTCCTCAGTGTTCCAGATGTCATCAACTGTCTGACAACAACTTATGATGG ACTTGAGCAAATGCATAAGGACCTGGTCAACGTTCCACTCTGTGTTGATATGTGT CTCAATTGGTTGCTCAATGTCTATGACACGGGTCGAACTGGAAAAATTAGAGTGC AGAGTCTGAAGATTGGATTAATGTCTCTCCCAAAGGTCTCTTGGAAGAAAAATA CAGATATCTCTTTAAGGAAGTTGCAGGGCCAACAGAAATGTGTGACCAGAGGCA GCTGGGCCTGTTACTTCATGATGCCATCCAGATCCCCGGCAGCTAGGTGAAGTA GCAGCTTTTGGAGGCAGTAATATTGAGCCTAGTGTTCGCAGCTGCTTCCAACAGA ATAACAATAAACCAGAAATAAGTGTGAAAGAGTTTATAGATTGGATGCATTTGG

AACCACAGTCCATGGTTTGGCTCCCAGTTTTACATCGAGTGGCAGCAGCGGAGAC TGCAAAACATCAGGCCAAATGCAACATCTGTAAAGAATGTCCAATTGTCGGGTTC AGGTATAGAAGCCTTAAGCATTTTAACTATGATGTCTGCCAGAGTTGTTTCTTTTC GGGTCGAACAGCAAAAGGTCACAAATTACATTACCCAATGGTGGAATATTGTAT 5 ACCTACAACATCTGGGGAAGATGTACGAGACTTCACAAAGGTACTTAAGAACAA CAGACAGTTCTTGAAGGTGACAACTTAGAGACTCCTATCACACTCATCAGTATGT GGCCAGAGCACTATGACCCCTCACAATCTCCTCAACTGTTTCATGATGACACCCA TTCAAGAATAGAACAATATGCCACACGACTGGCCCAGATGGAAAGGACTAATGG 10 GTCTTTCTCACTGATAGCAGCTCCACCACAGGAAGTGTGGAAGACGAGCACGCC CAGAGCCCAGCTCAGATCCTGAAGTCAGTAGAGAGGGAAGAACGTGGAGAACTG GAGAGGATCATTGCTGACCTGGAGGAAGAACAAAGAAATCTACAGGTGGAGTAT GAGCAGCTGAAGGACCAGCACCTCCGAAGGGGGCTCCCTGTCGGTTCACCGCCA 15 GAGTCGATTATATCTCCCCATCACACGTCTGAGGATTCAGAACTTATAGCAGAAG CAAAACTCCTCAGGCAGCACAAAGGTCGGCTGGAGGCTAGGATGCAGATTTTAG AAGATCACAATAAACAGCTGGAGTCTCAGCTCCACCGCCTCCGACAGCTGCTGG AGCAGCCTGAATCTGATTCCCGAATCAATGGTGTTTCCCCATGGGCTTCTCCTCA GCATTCTGCACTGAGCTACTCGCTTGATCCAGATGCCTCCGGCCCACAGTTCCAC 20 CAGGCAGCGGAGAGCCTGCTGGCCCCACCGCACGACACCAGCACGGATCTC ACGGAGGTCATGGAGCAGATTCACAGCACGTTTCCATCTTGCTGCCCAAATGTTC CTACAGTGTTGCCCTTTTCAGCAAATGCCAATTCCAAGTTCCATTAAATCAGAAG CECCATGGETCETTGGCCCACGATGTTGAGTGCTGACTGTGTGTTGTACTGAAAG::: 25 Á ÁGTÁAAACACTGACTATCCAAAGAGAAATGGATATTTTGTTTTTATAATAACCAT ATATTATTGTTTCTTCTCCCTTTCTATGCAAGTGTAAATTAATGAACAGAGAGG TATTTGGAAATGGTAATACATTTGTCACGGATTTGTATAATGTATACAGCATTGG GAAAGTGGGTGGGGCTTTCTAATATGATACCGTCTTTTTAATAACTATGACAAA GCTTACATAAGAATTAGAAGACCACTTTACATTTTTACATTCCTTCTGCTGTTCAT 30 ATTAACCTTGCACAATTACTTCATTTTTTTTTTTGACTCTTTTACCACAATGTTTTGG TTATTATAATTTATCAGCCATATGTTTATCAGCCATATAACCAACTAGATCCCAA ATAGATCCATGTATTTGTTTCCGTGATTTGGCCACATTAATAAATTCATAAATTTC AATCAAATATCATATATACACACATATGGTTTAAGCTACAGCCCTGTGTATGC CGTTTAACTTTATTTGACGTTGCCCACTTACTTCTTTGCTGACCACTTGGATAACC 35 GTAATAAAAATCCTATAAGCCTAAATGGCATTTCTTTTGGGATATTTTTCCTGCAT TGATAAAGAAGACTACATTATAATAATCTCAAAGATCATATTACCAAAGGTTGCC CACTTGAGCATATTTCATTTTGACACAGAAACAAAATTTAGTACAACCTTTCCT AGTTCCCATGTCTTGATTTCATCATTACATGCACAGCAGACCTTTACCTATTGTG 40 ATACCAGAACACATCATTGTCTTTGGTTCCCTTCAAAGAGAATTTTATTGTTGTTT TGTATTTCAAGTCCTTAATAGTTCTTGAAACTCCTAGTTGTTTCTTGTTGAAAG CAGACACACATTTAGTGCACGGCTTATTTTACCTTTCGGGTGAAAGATCAGATGT TTTTATACCCTTCACTTGATCAATATATTTGGAAAGAATGTTTATCAAAAGTCTAT GTCACTGCTTCTACAGAAGAATGAAATTAATGCTTAGGTGATGGTACCTCCACCT 45 ACATCTTTTGAGTGCATTCAATTATGTATTTTGGTTTAGCTTCTGATTTAACATTT AATTGATTCAGTTTAAACATGTTACTTAATTAGCAAATGTAGAGGAACCAAAAAA AGGTGAAAATAATATGTTTTGATTCAAACCTAAAGACATAAAAACATAAAGACA TTTTAACTTTGGGTTCTCTTTAGCTGGGATCTGGCCAGAAGGAGGCTTAAAGTTA GAAATTGCTATTATTTTAGAATAGGTTGGGTGGGTTGGGGGGCAAGGGTGTCTAT

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GAAGCCGAGCGTCTGAGCCTTCTGTGGGGCCGGTGGGGTTCTCACTGCGCTGGC AGCAGAGGATCTGCCTAAAGGTGGCGCTCATTTCTTTGTCGCGGTAGGAGTAAAT GGCCAGCACGTCGCACTGTGGACAGCACACGTCTAGAAGTAACAAAACCAATCC 5 AGGAGTCCAGCAGATGATAAAGGCCCCAAGCACAATGACCACAGTCTTCAGAAG ACTCATCATGGTATCCCGATTCCGCCGGGGTCCAGAACTATGCCGAGACATTCTC ATAGTCCTCTGGCGAACATAGCCAAAGATGTGAGCATAGAGAACCACCATTACC AGGGGTGCCATGTTGGAACAATTTTCAATATCACAGATACAGTTCCAGCCCACAC 10 TGGGTATAGCACCCATAACGATGGCCATAGTCCAGATGACCACAATGACCACCA CTACCGCCGGTTGCTCATCCGTGTGTGGAGCTGCATGCGGAAAACCGTAATGTG CCTCTCGATTGCAATAGCCAGTAAGTTGGCCACAGATGCCGTCAGGCTGGTGTCA CCTGTGTTGAACATGAGATAGAAGTAGGCCAACCCAGCAAAGAAGTCTGCAGCA 15 GCCAGATTAGCCATTAGGTAATAAATAGGAAAATGGAAGCGGCGGTTGACATAG ATTGCCACCATGACCAATAGGTTGGCCAACATGATGAAGATACAAACAGTGATT CCAAGTCCCATCACCAGCTTGCTGACTGTGTTCCATTCTGTGGCAAGATGCTTTCC ACTTCGGTTATAAAAGAAGGCAATGGACTCGTTGTAGAAGCACTGTGGTTCATTC ATGGCTGTGAACTGGGGCTGTGAAATTACAGGGATGGAAGTAGAGATGGCAGCC 20 ATGACAGCTCTGTGGTTGTAGGTGGTGAACACGCCCCAGAACTACGGGAGACAA ATTTTCTTGTTTGCTGATCAGATCGAAGTCATGCTAGGAGAAGCTGTGTACCTGA AND AND ATTEMPORATE AND ACCIDENCE OF THE STATE OF THE STA 😘 📑 CAGAATCCATGETGAGTGCCCACAGACCTGGGCAGGAGCTGTTCCCCCAGCGCCG 💠 25 GACAGCTGGCAGGACTCCGGTGGACGCCCCGGCACGGGCATTTTCACGTTGTC GCTCTCCTCTTCCCACTTGAAAAGCTCTGGAAAACATCGCGGGGCCCGCAAAACC CCGGAAATGTGGC

SEQ ID NO: 668

>21707 BLOOD 1147849.1 J03004 g183181 Human guanine nucleotide-binding regulatory protein (G) alpha-inhibitory-subunit mRNA, complete cds. 5e-78
 GCTGCACCGTGAGCGCCGAGGACAAGGCGGCGGCGGCGGGGAGGTGAAGTTGCTG
 ACAAGAACCTGCGGGAGGACGAGAGAAGGCGCGCGGGGAGGTGAAGTTGCTG
 CTGTTGGGTGCTGGGGAGTCAGGGAAGAGCACCATCGTNAAGCAGGTTAGGTCA
 TTNCCGGGGTTGTTATTTCCGGGGGGATTTCCNCAATACCCNGGGTTNTCTACAG
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 GACTTTGCCGACCCCTCC

SEQ ID NO: 669

>25177 BLOOD Hs.227948 gnl|UG|Hs#S553844 squamous cell carcinoma antigen=serine protease inhibitor [human, mRNA, 1711 nt] /cds=(61,1233) /gb=S66896 /gi=239551 /ug=Hs.227948 /len=1711
 CTCTCTGCCCACCTCTGCTTCCTCTAGGAACACAGGAGTTCCAGATCACATCGAG TTCACCATGAATTCACTCAGTGAAGCCAACACCAAGTTCATGTTCGACCTGTTCC
 45 AACAGTTCAGAAAAATCAAAAGAGAACAACATCTTCTATTCCCCTATCAGCATCAC ATCAGCATTAGGGATGGTCCTCTTAGGAGCCAAAGACAACACTGCACAACAGAT TAAGAAGGTTCTTCACTTTGATCAAGTCACAGAGAACACCACAGGAAAAGCTGC AACATATCATGTTGATAGGTCAGGAAAATGTTCATCACCAGTTTCAAAAGCTTCTG ACTGAATTCAACAAATCCACTGATGCATATGAGCTGAAGATCGCCAACAAGCTCT

TCGGAGAAAAACGTATCTATTTTACAGGAATATTTAGATGCCATCAAGAAATT TTACCAGACCAGTGTGGAATCTGTTGATTTTGCAAATGCTCCAGAAGAAGTCGA AAGAAGATTAACTCCTGGGTGGAAAGTCAAACGAATGAAAAAATTAAAAACCTA ATTCCTGAAGGTAATATTGGCAGCAATACCACATTGGTTCTTGTGAACGCAATCT 5 ATTTCAAAGGCCAGTGGGAGAAGAAATTTAATAAAGAAGATACTAAAGAGGAA AAATTTTGGCCAAACAAGAATACATACAAGTCCATACAGATGATGAGGCAATAC ACATCTTTCATTTTGCCTCGCTGGAGGATGTACAGGCCAAGGTCCTGGAAATAC CATACAAAGGCAAAGATCTAAGCATGATTGTTGCTGCCAAATGAAATCGATG GTCTCCAGAAGCTTGAAGAGAAACTCACTGCTGAGAAATTGATGGAATGGACAA 10 GTTTGCAGAATATGAGAGAGACACGTGTCGATTTACACTTACCTCGGTTCAAAGT GGAAGAGAGCTATGACCTCAAGGACACGTTGAGAACCATGGGAATGGTGGATAT CTTCAATGGGGATGCAGACCTCTCAGGCATGACCGGGAGCCGCGGTCTCGTGCTA TCTGGAGTCCTACACAGGCCTTTGTGGAGGTTACAGAGGAGGAGCAGAAGCT GCAGCTGCCACCGCTGTAGTAGGATTCGGATCACCTGCTTCAACTAATGAAG 15 AGTTCCATTGTAATCACCCTTTCCTATTCTTCATAAGGCAAAATAAGACCAACAG CATCCTCTTCTATGGCAGATTCTCATCCCCGTAGATGCAATTAGTCTGTCACTCCA TTTGGAAAATGTTCACCTGCAGATGTTCTGGTAAACTGATTGCTGGCAACAACAG ATTCTCTTGGCTCATATTTCTTTTCTTTCTCATCTTGATGATGATCGTCATCATCAA GAATTTAATGATTAAAATAGCATGCCTTTCTCTCTTTTCTCTTAATAAGCCCACATA 20 TAAATGTACTTTTCCTCCAGAAAAATTCTCCTTGAGGAAAAATGTCCAAAATAA GATGAATCACTTAATACCGTATCTTCTAAATTTGAAATATAATTCTGTTTGTGACC AND THE TETAL AN A STANCACACACATTECTTEGAATTTAGGTGATACCTAAATCCTTCTTATGTTCTAAATE . C. M. C. TTGTGATTCTATAAAACACATCATCAATAAAATAGTGACATAAAATCATAAAACACATCATCATAAAATAAAATAGTGACATAAAATCATAAAAACACATCATCAATAAAATAGTGACATAAAAATCATAAAAACACATCATCAATAAAATAGTGACATAAAAATCATAAAAATAA 25 AAAAAAAAAA

SEQ ID NO: 670

yc03e09.s1 Stratagene lung (#937210) Homo sapiens cDNA clone IMAGE:79624 3', mRNA sequence gi|666284|gb|T62627.1|T62627[666284]

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SEQ ID NO: 671

ys88a08.s1 Soares retina N2b5HR Homo sapiens cDNA clone IMAGE:221846 3' similar to SP:HTLF_HUMAN P32314 HUMAN T-CELL LEUKEMIA VIRUS ENHANCER FACTOR; contains MER22 repetitive element;, mRNA sequence

45 gi|1064703|gb|H84982.1|H84982[1064703]
GCTCCCAGTGGTCAGCGGAGACCCCAAGGAGGATCACAACTACAGCAGTGCCA
AGTCCTCCAACGCCCGGAGCACCTCGCCCACCAGCGACTCCATCTCCTCCTC
CTCCTCAGCCGACGACCACTATGAGTTTGCCACCAAGGGGAGCCAGGAGGCAG
CGAGGGCAGCGAGGGGAGCTTCCGGAGCCACGAGAGCCCCAGCGACACGGAAG

AGGACGACAGGAAGNACAGCCAGAAGGAGCCCAAGGATTTTTTNGGGGACAGC GGGTACGATTNCC

SEQ ID NO: 672

- 5 yq55b04.r1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:199663 5' similar to SP:SISD_HUMAN P13501 T-CELL SPECIFIC RANTES PROTEIN PRECURSOR; mRNA sequence gi|982328|gb|R96668.1|R96668[982328] NCGCCAGGAGTCCTCGGCCAGCCCTGCCTGCCCACCAGGAGATGAAGGTCTC CGTGGCTGCCCTCTCCTGCCTCATGCTTGTTGCTGTCCTTGGATCCCAGGCCCAGT
 10 TCACAAATGATGCAGAGACAGAGTTAATGATGTCAAAGCTTCCACTGGAAAATC
- 10 TCACAAATGATGCAGAGACAGAGTTAATGATGTCAAAGCTTCCACTGGAAAATC CAGTAGTTCTGAACAGCTTCACTTTGCTGCTGACTGCTGCACCTCCTACATCTCA CAAAGCATCCCGTGTTCACTCATGAAAAGTTATTTTGAAACGAGCAGCGAGTGCT CCAAGCCAGGGTGTCATATTCCTCACCAAGAAGGGGGCGGCAAGTCTGTGCCAAA CCCAGTGGGTCCGGGAGTTCAGGATTGGCATGGAAAAAGCTTNAAGCCCTAATT
- 15 CAATATTANTAATTAAAGGAGGACANAAGAGGGCCAGCNCACCCACCTCCAACA CTTCNTGAGGCTTTGGAAGG

SEQ ID NO: 673

- zt20b07.s1 Soares ovary tumor NbHOT Homo sapiens cDNA clone IMAGE:713653 3' similar to TR:G577291 G577291 MRNA ;contains element MER28 repetitive element ;,
- mRNA sequence
- - AGACAGAGGAAATTACGAGCCTCTGGGAGGGCAGCCTTTTCAATGCCAACTATG
 ACGTCCAGAGGTTCATTGTGGGATCAGACCGTGCTATCTTCATGCTTCGCGATGG
 GAGCTACGCCTGGGAGATCAAGGACTTTTTGGTCGGTCAAGACAGGTGTGCTGAT
 GTAACTCTGGAGGGCCAGGTGTACCCCGGCCAA GGAGGAGGAA

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SEQ ID NO: 674

>L01639

CGCATCTGGAGAACCAGCGGTTACCATGGAGGGGATCAGTATATACACTTCAGA TAACTACACCGAGGAAATGGGCTCAGGGGACTATGACTCCATGAAGGAACCCTG TTTCCGTGAAGAAAATGCTAATTTCAATAAAATCTTCCTGCCCACCATCTACTCC

- 35 TTTCCGTGAAGAAAATGCTAATTTCAATAAAATCTTCCTGCCCACCATCTACTCC
 ATCATCTTCTTAACTGGCATTGTGGGCAATGGATTGGTCATCCTGGTCATGGGTT
 ACCAGAAGAAACTGAGAAGCATGACGGACAAGTACAGGCTGCACCTGTCAGTGG
 CCGACCTCCTCTTTGTCACACGCTTCCCTTCTGGGCAGTTGATGCCGTGGCAAACT
 GGTACTTTGGGAACTTCCTATGCAAGGCAGTCCATGTCATCTACACAGTCAACCT
- 45 TATTGTCATCCTGCTATTGCATTATCATCTCCAAGCTGTCACACTCCAAGG GCCACCAGAAGCGCAAGGCCCTCAAGACCACAGTCATCCTCATCCTGGCTTTCTT CGCCTGTTGGCTGCCTTACTACATTGGGATCAGCATCGACTCCTTCATCCTCCTGG AAATCATCAAGCAAGGGTGTGAGTTTGAGAACACTGTGCACAAGTGGATTTCCA TCACCGAGGCCCTAGCTTTCTTCCACTGTTGTCTGAACCCCATCCTCTATGCTTTC

SEQ ID NO: 675

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> Human tumor necrosis factor receptor 2 (TNFR2) gene, exon 10 and complete cds 10 gi|1469539|gb|U52165.1|HSTNFR2S10[1469539] TCTTGGTCTCGGCTCTGGCCCAGTGCTCTTTCCCATGTGTCTGAATCTGCATCTT GGGCAGGGGTCCCTGGGCCCCACTCCTGGACCCCGGACTGACCCCCACCCCATC TTGTGCTTAGCAGATTCTTCCCCTGGTGGCCATGGGACCCAGGTCAATGTCACCT GCATCGTGAACGTCTGTAGCAGCTCTGACCACAGCTCACAGTGCTCCTCCCAAGC 15 CAGCTCCACAATGGGAGACACAGATTCCAGCCCCTCGGAGTCCCCGAAGGACGA GCAGGTCCCCTTCTCCAAGGAGGAATGTGCCTTTCGGTCACAGCTGGAGACGCCA GAGACCCTGCTGGGGAGCACCGAAGAGAAGCCCCTGCCCCTTGGAGTGCCTGAT GCTGGGATGAAGCCAGTTAACCAGGCCGGTGTGGGCTGTGTCGTAGCCAAGGT GGGCTGAGCCCTGCAGGATGACCCTGCGAAGGGGCCCTGGTCCTTCCAGGCCC 20 CCACCACTAGGACTCTGAGGCTCTTTCTGGGCCAAGTTCCTCTAGTGCCCTCCAC AGCCGCAGCCTCCCTCTGACCTGCAGGCCAAGAGCAGAGGCAGCGAGTTGGGGA 25 TTTGTTTCTCCCCTGGGCTCTGCCCAGCTCTGGCTTCCAGAAAACCCCAGCATCC TTTTCTGCAGAGGGCTTTCTGGAGAGGGAGGGATGCTGCCTGAGTCACCCATGAA GACAGGACAGTGCTTCAGCCTGAGGCTGAGACTGCGGGATGGTCCTGGGGCTCT GTGTAGGGAGGAGGTGGCAGCCCTGTAGGGAACGGGGTCCTTCAAGTTAGCTCA GGAGGCTTGGAAAGCATCACCTCAGGCCAGGTGCAGTGGCTCACGCCTATGATC 30 CCAGCACTTTGGGAGGCTGAGGCGGGTGGATCACCTGAGGTTAGGAGTTCGAGA CCAGCCTGGCCAACATGGTAAAACCCCATCTCTACTAAAAATACAGAAATTAGC CGGGCGTGGTGGCGGCACCTATAGTCCCAGCTACTCAGAAGCCTGAGGCTGGG AAATCGTTTGAACCCGGGAAGCGGAGGTTGCAGGGAGCCGAGATCACGCCACTG 35 GCACCGCCTCCAAATGCTAACTTGTCCTTTTGTACCATGGTGTGAAAGTCAGATG CCCAGAGGCCCAGGCAGCCACCATATTCAGTGCTGTGGCCTGGGCAAGATAA CAACAAGCCAACGACAAACTCTGCCAGCCACATCCAACCCCCACCTG 40 TGCTGTCCTAGGCCACACCATCTCCTTTCAGGGAATTTCAGGAACTAGAGATGAC TGAGTCCTCGTAGCCATCTCTCTACTCCTACCTCAGCCTAGACCCTCCTCCTCCCC CAGAGGGGTGGGTTCCTCTTCCCCACTCCCACCTTCAATTCCTGGGCCCCAAAC GGGCTGCCCTCCACTTTGGTACATGGCCAGTGTGATCCCAAGTGCCAGTCTTGT GTCTGCGTCTGTGCGTGTCGTGGGTGTGTAGCCAAGGTCGGTAAGTTGAA 45 TGGCCTGCCTTGAAGCCACTGAAGCTGGGATTCCTCCCCATTAGAGTCAGCCTTC CCCCTCCAGGGCCAGGGCCCTGCAGAGGGGAAACCAGTGTAGCCTTGCCCGGA TTCTGGGAGGAAGCAGGTTGAGGGGCTCCTGGAAAGGCTCAGTCTCAGGAGCAT

AATTGTTGATAAATTCCACTGGACTTGAGCTTGGCAGCTGAACTATTGGAGGGTG

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SEQ ID NO: SEQ ID NO: 676

>R88734

20 SEQ ID NO: 677 >AA418689

- 25 TCTTTCGGGCCTTGAGTTCCTTCATGGCAATGAGCAGAGGATCTGTCTCCCCCTCC
 AGCTCCACCATCACAGGGGCACACATCGCAATCTGGAGCGCTCGGGTGCCCAGC
 ACGCGGGCTCGCTCGTACTTGGTCATGTATGGTGTGATTCGCTTCTGGTTGG
 CCTGCGGTCGCTCCCCAGAGGAGGNAGTCTCGACATTCTCCTGGCCTTCCTCTC
 GGCATTCTCCAAGTCATCTAGCCCTTCATCCTCCTC
- 30 CACATCATCAGAGTCGTCGCCATCAAA

SEQ ID NO: 678 >AA455281

TTTTTGGAGGAGTGGCATGGAGTTCTTTAATTTGGAAGGCAAAAGGTTACATTTA

ATGAAAGGCAGAGGCTGGATTAATAAATGTTTGTTAGAAAGTTGTTCTGACACAC

AGTGAACTCTGGGCTTTTCTCCTGCATAAAAAGCAGAGCTAGCAGTAAGTGCAA

ATCTGAAGAAAATCCATGTGTCCAATAAGCTGCCATCTCCAGAACTCTTATCCAG

GAAATTCAAAGAGTGAACATTCTTTTAGTCTCCTACTCCTCAATTAAGTAAATGA

GAATGAGTCAGCCAACAAAGTTCATGACAACAAGGTGCAGGATGGTGCTGGCAA

40 AGAGAAAATCAGCAAAGGCTCGCTCTGGGGAGATGCCTTGGAAATCCGCTTTGT TCTGTGGGTTGATCTGTATTCTCAGGCAAACCGCTAGGATGAAACTCCCCACACA AGAGATGAAGCCCGAGAGAAAAGAGTTGAAGGGGAAGGTCCC

SEQ ID NO: 679

45 >H94469

GCAAAACAACATTTATTCTTTTAAAAAATCTATATACATTGCCATACAAAGATAC CACATTGAAGCAGTTCTCAGGAACCTTCCAGTGAGCCTTCTTATAATTGCCCG AGCAAGATTTCGTGCCAGAGAAAGTCTCAGCATTTCCACCTTGGTGTNCTCTATG TCATCATCCTGGAGCTGCTCGGTATCAGATTCTCCATGCACAGGTCTTCTTGACGT

CAAGTCCTCCAGACACCGCATCAACTCATAAGTCTGTTCTGCTGAGAAAATCACC TGTTTCTGTTCCAAAAGGGGCAAGGCATCTGTCAGCAGAGTTCATCCCAGAAAGA CCGAAGGGGCAATCCGAGACGTCATCAAG GACAGAAGGA

- 5 SEQ ID NO: 680
 - aa79c05.s1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:827144 3' similar to SW:RLX1_HUMAN P49406 PUTATIVE 60S RIBOSOMAL PROTEIN;, mRNA sequence gi|2261786|gb|AA521243.1|AA521243[2261786] TTTTTTTTTGGTGTACAAGTTTTATTTTAGAAAAAAAGTATTAATAAAACAATGA
- 15 AAATTTGGACGTTCCCAGCGTTTAGACCAGGGCTTAGGCTTCATTTTACTTTCAG CTCATTAACAGGAACTTTTTGGTTAGGCTCTTGTACTACTGGCTTCATATTCACAT CAAAAGTGCTATATTCAGGAAGGGCATCTCGTAAGTATAGCAAGCTATCATCCA GCCGTTTCTCTAATTTGACCACCTGAATCTCCTGGACCCGAGGATTATAAAGTTC AAAGCAAATCTCGACACCTTGTCCTTCGATAACATTCCTAAGGAT GAAAGTAGC

20 SEO ID NO: 681

- - GGATCCAGGACTGAGATCCCAGAACCATGAACCTGGCCATCAGGATCGCTCTCCT
 GCTAACAGGTACCCGGCATGGGGCAGGACTGGGCTCCAGGCGCCCTGGCTTCC
 - TTCCTCCAGAGAAGCAGCTTCTCCCTCACAGTCTCAGAAAAGCGCAGGTGACAA
 AGAGAGGGCTCTTTTTCATCCTGAAGTCAGCCGATCCACCGCGCTGATATTCTGA
 CGGCCTGAGGTGGTTTTTGGAAACACAGTTTGCTGAGCCCTCCTTCACACTATTG
 AACTAGAATCCCCAACTGAGAACCCAGGAACCAGCATCAACTCCCTAAGATCTC

 - 35 GTGGACCAGAGCCTTCGTCTGGACTGCCGCCATGAGAATACCAGCAGTTCACCCA TCCAGTACGAGTTCAGCCTGACCCGTGAGACAAAGAAGCACGTGCTCTTTGGCAC TGTGGGGGTGCCTGAGCACACATACCGCTCCCGAACCAACTTCACCAGCAAATA CCACATGAAGGTCCTCTACTTATCCGCCTTCACTAGCAAGGACGAGGGCACCTAC ACGTGTGCACTCCACCACTCTGGCCATTCCCCACCCATCTCCCCAGAACGTCA

 - 45 AACAGGATGACACCACCTCCCTCAGCCAGTTTTCTTGTCATGATGTTTAGTAAG
 GTTTTCATAAGATGATATGTGTGCAAGAGATCAGTAATCTGCAAATGGGAAAGA
 TGGCTGGTTCTGTGAGACCAGGCTGTTCCTGGTCCCAGCTAAGACATTGCAGTAC
 CCACCTCCCAAAGGGAGTACACCCTTGCTTTGGGCCTGTGCCTGAGTCCTG
 ATCCGTCTTCCTTCCTACCCTGCCCCCGGCCCCCTTCTCTTTCTGCAGACAAACTG

GTCAAGTGTGAGGGCATCAGCCTGCTGGCTCAGAACACCTCGTGGCTGCTGCTGC TCCTGCTGTCCCTCCCTCCAGGCCACGGATTTCATGTCCCTGTGACTGGTG GGGCCCATGGAGGAGACAGGAAGCCTCAAGTTCCAGTGCAGAGATCCTACTTCT CTGAGTCAGCTGACCCCCTCCCCCAATCCCTCAAACCTTGAGGAGAAGTGGGGA 5 CCCCACCCTCATCAGGAGTTCCAGTGCTGCATGCGATTATCTACCCACGTCCAC GCGGCCACCTCACCCTCTCGCACACCTCTGGCTGTCTTTTTGTACTTTTTGTTCC TGAAGAGGGAAGCCAGGATTGGGGACCTGATGGAGAGTGAGAGCATGTGAGGG GTAGTGGGATGGTGGGTACCAGCCACTGGAGGGGTCATCCTTGCCCATCGGGA 10 CCAGAAACCTGGGAGAGACTTGGATGAGGAGTGGTTGGGCTGTGCTGGGCCTAG GACCCCAGATGTGAGGGCACCACCAAGAATTTGTGGCCTACCTTGTGAGGGAGA GCCCTCCTTACCACTGTGGAAGTCCCTCAGAGGCCTTGGGGCATGACCCAGTGAA 15 GATGCAGGTTTGACCAGGAAAGCAGCGCTAGTGGAGGGGTTGGAGAAGGAGGTA AAGGATGAGGGTTCATCATCCCTCCCTGCCTAAGGAAGCTAAAAGCATGGCCCT GCTGCCCCCCCCCCCCCCCCCCACAGTGGAGAGGCTACAAAGGAGGACAAGA CCCTCTCAGGCTGTCCCAAGCTCCCAAGAGCTTCCAGAGCTCTGACCCACAGCCT CCAAGTCAGGTGGGGTGGAGTCCCAGAGCTGCACAGGGTTTGGCCCAAGTTTCT 20 TGAGCCCTCAGACAGCCCCTGCCCGCAGGCCTGCCTTCTCAGGGACTTCTGC with it GGGGCCTGAGGCAAGCCATGGAGTGAGACCCAGGAGCCGGACACTTCTCAGGAA ATAAAACCAAGCCTCTGGAATCTGTCCTCGTGTCCACCTGGCCTTCGCTCCTCCA 25 GCAGTGCCTGCCTGCCCCGCTT

SEQ ID NO: 682

yw08h11.s1 Soares melanocyte 2NbHM Homo sapiens cDNA clone IMAGE:251685 3',

30 mRNA sequence gi|1110224|gb|H96738.1|H96738[1110224]
TAAAANAAATCTTTTTTTATTTCAAAGATTGCTTCTTATATTGAAGCTCATATTA
AAGCAACAGTACAATGTTCATAAAATATAAGTGTGATGCCGTAACATTTTCTTAC
ATGTCAGAATACTGATATTTATATGTATACTAAAATAAGAACTTTAAAATTGTAC
AAATAGATACATTAAAAATGACATAGAAATAGGGCGTCTCTCACTGAAACAAGA

35 CAGTTATATCTGGCACGTATTAGTTTAAGATGAAAGTAGAAGCAAAAAGATTTAC AAGAATCAGCAGTAACAAGATTGATGCTCAAGAGACATAATTGTACATTGTATT GTACATACATTGTATGGGTTTAAGCTGGCTGGAATATTATATATTTCCAAGTTTTA AAAATGGCNCTACCANATAGAGTGGTCCNGAGTTTAAGGCGAAATTACAGCTCA GAACTGTTGTCCCTTCNAATTTTGGTGG

SEO ID NO: 683

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Human integral membrane serine protease Seprase mRNA, complete cds gi|1924981|gb|U76833.1|HSU76833[1924981]

CCACGCTCTGAAGACAGAATTAGCTAACTTTCAAAAAACATCTGGAAAAATGAAG

45 ACTTGGGTAAAAATCGTATTTGGAGTTGCCACCTCTGCTGTGCTTATTGGT
GATGTGCATTGTCTTACGCCCTTCAAGAGTTCATAACTCTGAAGAAAATACAATG
AGAGCACTCACACTGAAGGATATTTTAAATGGAACATTTTCTTATAAAACATTTT
TTCCAAACTGGATTTCAGGACAAGAATATCTTCATCAATCTGCAGATAACAATAT
AGTACTTTATAATATTGAAACAGGGCAATCATATACCATTTTGAGTAATAGAACC

ATGAAAAGTGTGAATGCTTCAAATTACGGCTTATCACCTGATCGGCAATTTGTAT ATCTAGAAAGTGATTATTCAAAGCTTTGGAGATACTCTTACACAGCAACATATTA CATCTATGACCTTAGCAATGGAGAATTTGTAAGAGGAAATGAGCTTCCTCGTCCA ATTCAGTATTTATGCTGGTCGCCTGTTGGGAGTAAATTAGCATATGTCTATCAAA 5 ACAATATCTATTTGAAACAAGACCAGGAGATCCACCTTTTCAAATAACATTTAA TGGAAGAGAAAATATATTAATGGAATCCCAGACTGGGTTTATGAAGAGGA AATGCTTGCTACAAAATATGCTCTCTGGTGGTCTCCTAATGGAAAATTTTTGGCA TATGCGGAATTTAATGATACGGATATACCAGTTATTGCCTATTCCTATTATGGCG ATGAACAATATCCTAGAACAATAAATATTCCATACCCAAAGGCTGGAGCTAAGA 10 ATCCCGTTGTTCGGATATTTATCGATACCACTTACCCTGCGTATGTAGGTCCC CAGGAAGTGCCTGTTCCAGCAATGATAGCCTCAAGTGATTATTATTTCAGTTGGC TCACGTGGGTTACTGATGAACGAGTATGTTTGCAGTGGCTAAAAAGAGTCCAGA ATGTTTCGGTCCTGTCTATATGTGACTTCAGGGAAGACTGGCAGACATGGGATTG TCCAAAGACCCAGGAGCATATAGAAGAAAGCAGAACTGGATGGGCTGGTGGATT 15 CTTTGTTTCAACACCAGTTTTCAGCTATGATGCCATTTCGTACTACAAAATATTTA GTGACAAGGATGGCTACAAACATATTCACTATATCAAAGACACTGTGGAAAATG CTATTCAAATTACAAGTGGCAAGTGGGAGGCCATAAATATATTCAGAGTAACAC AGGATTCACTGTTTTATTCTAGCAATGAATTTGAAGAATACCCTGGAAGAAGAAA 20 CATCTAAGGAAAGAAAGGTGCCAATATTACACAGCAAGTTTCAGCGACTACGCC AAGTACTATGCACTTGTCTGCTACGGCCCAGGCATCCCCATTTCCACCCTTCATG **XTGCTTTGAAAAATATCCAGCTGCCTAAAGAGGAAATTAAGAAACTTGAAGTAG ATGAAATTACTTTATGGTACAAGATGATTCTTCCTCCTCAATTTGACAGATCAAA 25 GAAGTATCCCTTGCTAATTCAAGTGTATGGTGGTC@CTGCAGTCAGAGTGTAAGG TTGCCTTGGTGGATGGTCGAGGAACAGCTTTCCAAGGTGACAAACTCCTCTATGC AGTGTATCGAAAGCTGGGTGTTTATGAAGTTGAAGACCAGATTACAGCTGTCAG AAAATTCATAGAAATGGGTTTCATTGATGAAAAAAGAATAGCCATATGGGGCTG 30 GTCCTATGGAGGATACGTTTCATCACTGGCCCTTGCATCTGGAACTGGTCTTTTCA CACAGAGAGATTCATGGGTCTCCCAACAAGGATGATAATCTTGAGCACTATAA GAATTCAACTGTGATGGCAAGAGCAGAATATTTCAGAAATGTAGACTATCTTCTC ATCCACGGAACAGCAGATGATAATGTGCACTTTCAGAACTCAGCACAGATTGCT 35 AAAGCTCTGGTTAATGCACAAGTGGATTTCCAGGCAATGTGGTACTCTGACCAGA ACCACGGCTTATCCGGCCTGTCCACGAACCACTTATACACCCACATGACCCACTT CCTAAAGCAGTGTTTCTCTTTGTCAGACTAAAAACGATGCAGATGCAAGCCTGTA **TCAGAATCTGA**

SEQ ID NO: 684
 zw83d07.s1 Soares_testis_NHT Homo sapiens cDNA clone IMAGE:782797 3', mRNA sequence gi|2161864|gb|AA448194.1|AA448194[2161864]
 TTTTTTTTAAAAAAAAAATTAAATATTTTTATTATATACTTTTAAACATATAGAAGA TAGAAAAAAAACAGTACAATGAACAGCCATGTCCACCAGTTAGATTCTGTAACAT
 TTTGCCACATACGCCTCACATACATTTTGTTAAACCATTTGAAACATTTTAAGACA CTCTAACACTTCATTCCTAAATGCTTAAGTATGCAAATTAAGACAGTCTTTTATAA ACTACAACACCCTTCTCACAGGCTCATAAAATTACCAATAATTATCCAATATCATT CAAAATCTAATCCACATTCAAATTTTCTCAACTGCCTCACCACCGTGCTGGCCTCC

CACCCCCACCTCAGTCTTTTACAGATGGTTTTTCAAAATAGAGTCCAGTAAAATA TTTCACATTGCATTTGGTTATTACATAACTTT TAATCAAGAAGAGTTAC

SEQ ID NO: 685

- 5 Human gene for preproenkephalin gi|31150|emb|V00509.1|HSENK1[31150] CCGACCCTCCCGCGAAGGCGTCGGCGCGGGGCTGGCGTAGGGCCTGCGTCAGC TGCAGCCGCCGCGATTGGGGCGCGCGCCCTCCTTCGGTTTGGGGCTAATTAT CCCGCAGCCTGGCCCGTGACCCCGCAGAGACGCTGAGGACCGCGACGGTGAGGC 10
- ACTTGCCTTCTTCCCTCTAGAGTCGTGTCTGAACCCGGCTTTTCCAATTGG CCTGCTCCATCCGAACAGCGTCAACGTGAGTGAATTTGCCCGAAGCTTGTCTTTG CTGAGCGGGTTTGGGGACGTCTGCCCGCCCTCTTTCCCTTCACATTTCATTGCATG GGTTCCCCAACAGCGTTCCCTGGTTCTTTTTGTGACCCCAGTCAATGTCCTGCCT
- 15 CCCCGGCTCCCGCTCTCTCGCCCCTGGTCTGCGGCGTTCTCTCCGGAATCTTGCC CTGGGCCGCGGACGCCCAGGAAAAGAGCCGGGTGCCCCAGGCAGCCTCGCGTTG GGGGCGACCGCCATCCCGGGAACCGCGAGGCGATCTGAGTCGCCTCCACGTC TACCTAAAAGCTGTCGGCCGGGAGGGCGGGGCCCCAGAAAGGAGCATTCCTGCG GGCTTTTGCTCGACGATCCCCTGCTGAGGCTGTCGCGGCGAGGGTCCTGCCGAGG
- 20 GACCCCGTTCTGCGCCCAGGCAGGCTCGAAGCACGCGTCCCTCTCTCCTCGCAGT CCATGGCGCGGTTCCTGACACTTTGCACTTGGCTGCTGTTGCTCGGCCCCGGGCT **********CETAGTGCGCCGGCCGACATCAACTTCCTGGTGAGTGTTGCGCGCGGGGAGTGT

LANDA & TGCGCACCTTGTGAGACAGAGTTTCCG SALES, Despite the first of the second state of th

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SEQ ID NO: 686

yi26g12.s1 Soares placenta Nb2HP Homo sapiens cDNA clone IMAGE:140422 3', mRNA sequence gi|838397|gb|R65759.1|R65759[838397]

- AAAATTTTTTTTACCGTATTTATTGGTTCAAAAACTAGAATTTATAGTTTCAGGCA 30 GATTTCAACCAAAGAGTCACCAAATTAAATACACAGGGTAGCTTGTGAGGCATA GACACAGCCCATGTGTTTTCCTCTACATTGTATATTCATTTCTCTTTTGGCGATTTG ACATTATAGCCATTCTCTGGAAGTCCTAAAGCAAACTAGTATTTTATGTGCCATA TTAAGTTAAAATTTCTTATGTGAGGATACCACTAATACTGGGTTTTGATTTAGGG CCATCCTTCTTGCCGGGGGTATGGACAATGGGGGGCTTGTTTCTATGGATTAAG 35 GNCCCTACCCCTGGGGCCAGGTGNTATGGGGGGNATTGTTAAAACCATGGCCATT
- ATTATGGTGGGGGCCAACCCCCACCCNTGGAAG GGGA

SEO ID NO: 687 >R91550

- 40 GGAGGATGTGGGCCACGCAGGCTGGCGGTGGCGCTGGCTCTGAGCGTGCCC GGGCACCGGGCGCCGGCCGGCGACTGCGAAGTTTGTATTTCTTATCTGGGAA GATTTTACCAGGACCTCAAAGACAGAGATGTCACATTCTCACCAGCCACTATTGA AAACGAACTTATAAAGTTCTGCCGGGAAGCAAGAGGCAAAGAGAATCGGTTGTG CTACTATATCGGGGCCACAGATGATGCAGCCACCAAAATCATCAATGAGGTATC
- 45 AAAGCCTCTGGCCCCACCACATCCCTGTGGGAGAAGATCTGTGAGAAGCTTAAG GAAGAAGGACAGCCAGATATGTGAGCTTAAGTAT GGACAAGCAGATCC

SEQ ID NO: 688 >M94054

GGGCGTGATTTGAGCCCCGTTTTTATTTTCTGTGAGCCACGTCCTCCTCGAGGGG GTCAATCTGGCCAAAAGGAGTGATGCGCTTCGCCTGGACCGTGCTCCTGGG CCTTTGCAGCTCTGCGCGCTAGTGCACTGCGCCCCTCCCGCCGCCGGCCAACAGC AGCCCCGCGCGAGCCGCCGGCGCTCCGGGCGCCTGGCGCCAGCAGATCCAAT 5 GGGAGAACAACGGCAGGTGTTCAGCTTGCTGAGCCTGGGCTCACAGTACCAGC CTCAGCGCCGCGGGACCCGGGCGCCGCCGTCCCTGGTGCAGCCAACGCCTCCG CCCAGCAGCCCGCACTCCGATCCTGCTGATCCGCGACAACCGCACCGCCGCGC GCGAACGCGGACGCCGCTCATCTGGAGTCACCGCTGGCCGCCCCAGGCCCAC CGCCCGTCACTGGTTCCAAGCTGGCTACTCGACATCTAGAGCCCGCGAACGTGGC 10 GCCTCGCGCGGAGAACCAGACAGCGCCGGGAGAAGTTCCTGCGCTCAGTAAC CTGCGGCCGCCAGCCGCGTGGACGCCATGGTGGGCGACGACCCTTACAACCCC TACAAGTACTCTGACGACAACCCTTATTACAACTACTACGATACTTATGAAAGGC CCAGACCTGGGGCAGGTACCGGCCCGGATACGGCACTGGCTACTTCCAGTACG 15 GAAGATGTCCATGTACAACCTGAGATGCGCGGCGGAGGAAAACTGTCTGGCCAG TACAGCATACAGGGCAGATGTCAGAGATTATGATCACAGGGTGCTGCTCAGATTT CCCCAAAGAGTGAAAAACCAAGGGACATCAGATTTCTTACCCAGCCGACCAAGA TATTCCTGGGAATGGCACAGTTGTCATCAACATTACCACAGTATGGATGAGTTTA GCCACTATGACCTGCTTGATGCCAACACCCAGAGGAGAGTGGCTGAAGGCCACA 20 AAGCAAGTTTCTGTCTTGAAGACACATCCTGTGACTATGGCTACCACAGGCGATT TGCATGTACTGCACACACACAGGGATTGAGTCCTGGCTGTTATGATACCTATGGT # GCAGACATAGACTGCCAGTGGATTGATATTACAGATGTAAAACCTGGAAACTAT ATCCTAAAGGTCAGTGTAAACCCCAGCTACCTGGTTCCTGAATCTGACTATACCA 25 CTGCACAATTTCACCGTATTAGAAGGCAAAGCAAAACTCCCAATGGATAAATCA GTGCCTGGTGTTCTGAAGTGGGAAAAAATAGACTAACTTCAGTAGGATTTATGTA TAACAAAGCACATAACTGGATTTTGAACGCTTAAGTCAATCATTACTTGGAAATT TNTAATGTTTATTATTACATCAACTTTGTGAATTAACACAGTGTTTCAATTCTGT 30 **AATTTCATATTTGACTCTTT**

SEO ID NO: 689

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Human mRNA for beta-actin gi|28251|emb|X00351.1|HSAC07[28251] TTGCCGATCCGCCGCCGTCCACACCCGCCGCCAGCTCACCATGGATGATAT CGCCGCGCTCGTCGACAACGCTCCGGCATGTGCAAGGCCGGCTTCGCGGG CGACGATGCCCCCGGGCCGTCTTCCCCTCCATCGTGGGGCGCCCCAGGCACCAG GGCGTGATGGTGGGCATGGGTCAGAAGGATTCCTATGTGGGCGACGAGGCCCAG AGCAAGAGAGGCATCCTCACCCTGAAGTACCCCATCGAGCACGGCATCGTCACC AACTGGGACGACATGGGAAAAATCTGGCACCACACCTTCTACAATGAGCTGCGT GTGGCTCCCGAGGAGCACCCCGTGCTGCTGACCGAGGCCCCCCTGAACCCCAAG GCCAACCGCGAGAGATGACCCAGATCATGTTTGAGACCTTCAACACCCCAGCC ATGTACGTTGCTATCCAGGCTGTGCTATCCCTGTACGCCTCTGGCCGTACCACTG GCATCGTGATGGACTCCGGTGACGGGGTCACCCACACTGTGCCCATCTACGAGG TGACTACCTCATGAAGATCCTCACCGAGCGCGGCTACAGCTTCACCACCACGGCC GAGCGGGAAATCGTGCGTGACATTAAGGAGAAGCTGTGCTACGTCGCCCTGGAC TTCGAGCAAGAGATGGCCACGGCTGCTTCCAGCTCCTCCCTGGAGAAGAGCTAC GAGCTGCCTGACGGCCAGGTCATCACCATTGGCAATGAGCGGTTCCGCTGCCCTG AGGCACTCTTCCAGCCTTCCTTGGGCATGGAGTCCTGTGGCATCCACGAAAC

TACCTTCAACTCCATCATGAAGTGTGACGTGGACATCCGCAAAGACCTGTACGCC AACACAGTGCTGTCTGGCGCACCACCATGTACCCTGGCATTGCCGACAGGATGC AGAAGGAGATCACTGCCCTGGCACCCAGCACAATGAAGATCAAGATCATTGCTC CTCCTGAGCGCAAGTACTCCGTGTGGATCGGCGGCTCCATCCTGGCCTCGCTGTC 5 CACCTTCCAGCAGATGTGGATCAGCAAGCAGGAGTATGACGAGTCCGGCCCCTC CATCGTCCACCGCAAATGCTTCTAGGCGGACTATGACTTAGTTGCGTTACACCCT TTCTTGACAAAACCTAACTTGCGCAGAAAACAAGATGAGATTGGCATGGCTTTAT TTGTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTAA 10 TCACAATGTGGCCGAGGACTTTGATTGCACATTGTTGTTTTTTTAATAGTCATTCC AAATATGAGATGCATTGTTACAGGAAGTCCCTTGCCATCCTAAAAGCCACCCCAC TTCTCTCTAAGGAGAATGGCCCAGTCCTCTCCCAAGTCCACACAGGGGAGGTGAT AGCATTGCTTTCGTGTAAATTATGTAATGCAAAATTTTTTTAATCTTCGCCTTAAT ACTTTTTATTTTGTTTTTTTGAATGATGAGCCTTCGTGCCCCCCTTCCCCCTT 15 AGGCAGCCAGGCTTACCTGTACACTGACTTGAGACCAGTTGAATAAAAGTGCA CACCTTA

SEQ ID NO: 690

20 >AA435938

TTTCATGCTCATTGCTGTTTATTGAAACAAAAGAATCAGAAGAAGATCAGAATGA GATATGTCATGGAAGGCTTCTTTAAACACCCAGAAGAAATTCAGGATAAAGCTCA AAAAGAGCAGGCAATCGATAGGGGTTGAAAATCCACTCAGTAGGCCACGGAAG GACTTCAAGAAGGTTGATCGTTCTGTCGCTGGATGTTGTAGGTGTCCTACGTGAA GGCAATCGACATCTGGATGGCTGTGTGTCTCTTTTGTGTT

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SEQ ID NO: 691 >AA443497

CGCTGCCTTGCTGGAG

TCCAAGGTCATGGCAAAACATCTGAAGTTCATCGCCAGGACTGTGATGGTACAG GAAGGGAACGTGGAAAGCGCATACAGGACCCTAAACAGAATCCTCACTATGGAT GGGCTCATTGAGGACATTAAGCATCGGCGGTATTATGAGAAGCCATGCCGCCGC 35 GACAGAGGGAAAGCTATGAAAGGTGCCGGCGGATCTACAACATGGAAATGGCTC GCAAGATCAACTTCTTGATGCGAAAGAATCGGGCAGATCCGTGGCAGGGCTGCT GAGGCCTGTGGGTGGGACACCAGTGCGAAACCCTCATCCAGTTTTCTCTCCATCT CTTTTCTTGTACAATCCCATTTCCTATTACCATTCTCTGCAATAAACTCAAATCA 40 **CATGTCTGC**

SEQ ID NO: 692 zf17e01.s1 Soares fetal heart NbHH19W Homo sapiens cDNA clone IMAGE:377208 3', mRNA sequence gi|1547536|gb|AA055198.1|AA055198[1547536]

45 CACCTTAAAAACTAGGTTTCTATTTCTGGTTAGATTCTAGAGCAGTGGAACTCAG AGATAACATTGTACAAAACTGTATTTACAAGAAAACCAATTAAAAATTAAGGGT GTGTGCAAAAGTAGACAGGAGAGTCAAGACATATCAATGCAGGGATGGCTTTGG GGAATGGGGACTCAAGGTTCTACACTGGAACCTGGGG

SEQ ID NO: 693

zt87h10.s1 Soares_testis_NHT Homo sapiens cDNA clone IMAGE:729379 3', mRNA sequence

- 5 gi|2140847|gb|AA435933.1|AA435933[2140847]
 - TTTTGGTTCAAACAATGGAACATTTTATTATTATCATATTACAAAGAGTCAGTGAT GGGCC
- 10 TATATCAAGATGCAGTATTCACAGAAAGAGGACTGTTCATTTCTTTACCAGAAGA TTCTCCCATATATCATGTGTCTACATCTAAACCAATCACTACTAAGGGGAAATTG ACCTACAACATTTGGATTAGACTAATCAAATTTACCTTCTGAGTTAGGCATAGAG TCAACTTCTATGAGCACATGGCTGAGCCAAGGATAAGCATTCTGCCAGCAAGAG AGGACATAATATGGGTGTGGGGATTGGAGATGGGAGAG

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- **SEQ ID NO: 694**
- yo27c07.s1 Soares adult brain N2b5HB55Y Homo sapiens cDNA clone IMAGE:179148 3', mRNA sequence gi|989944|gb|H50103.1|H50103[989944]
- AAATTTATCAATGACAAACAGACATAAAACTCAAAGTTTGGCTCTTCTGAGGGGC AGGAGAAAAACTGGTGATGTTCTTTTATACAGATGAAACATGGGTNCAGAAATT
- ** CAACGTCTCEAGCGCTTAGGNCCGTAAAANTGTTCTAAGCACAGAAGTACATGT
 - 25 GGGAAGATTTCTCTCATCATTTTTNGTAAANCAAAGCGTTCTAATATTTTACAGA CCAAGTTAGGGCCAGTTTTTNTTTTTCCCT

SEQ ID NO: 695

- za29f01.r1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:293977 5',
- 30 mRNA sequence gi|1267964|gb|N95657.1|N95657[1267964]
 - GCAGAAGCGAACAACCTGAGCTTTCCCTTGGAGCCCCTGAGCAGGGAGAGGGCT CACAAGCTTGAGGCCATCTCTCGCCTCTGCGAGNACNAAGTACAAGGACCTAAG AAGATCCGCGAGAAGCGCTCAGCCAGTGCAGACAACCTGACTCTGCCCCGGTGG TCCCCAGCCATCATCTCTTAACTACGGAGGCCCGCCGGACCACACCATCCCTTAG
- 35 TTTCTCCTTTAGTTTGAGAAAAGACAGACTTGGGGTNGGTTTGTTTTTTTC TTTCCTTTTTTTTACGCATAGCTCCCGTCAAAGCTGCCT

SEQ ID NO: 696

- Human lysophosphatidic acid receptor homolog mRNA, complete cds
- 40 gi|1857424|gb|U80811.1|HSU80811[1857424]
 - TCACCACCTACAACCACAGAGCTGTCATGGCTGCCATCTCTACTTCCATCCTGT AATTTCACAGCCCCAGTTCACAGCCATGAATGAACCACAGTGCTTCTACAACGAG TCCATTGCCTTCTTTTATAACCGAAGTGGAAAGCATCTTGCCACAGAATGGAACA CAGTCAGCAAGCTGGTGATGGGACTTGGAATCACTGTTTGTATCTTCATCATGTT

CGGATGAGCAACCGGCGGGTAGTGGTGGTCATTGTGGTCATCTGGACTATGGCC ATCGTTATGGGTGCTATACCCAGTGTGGGCTGGAACTGTATCTGTGATATTGAAA TTCAACTTGGTGACCTTTGTGGTAATGGTGGTTCTCTATGCTCACATCTTTGGCTA 5 TGTTCGCCAGAGGACTATGAGAATGTCTCGGCATAGTTCTGGACCCCGGCGGAAT CGGGATACCATGATGAGTCTTCTGAAGACTGTGGTCATTGTGCTTGGGGCCTTTA TCATCTGCTGGACTCCTGGATTGGTTTTGTTACTTCTAGACGTGTGCTGTCCACAG TGCGACGTGCTGGCCTATGAGAAATTCTTCCTTCTCCTTGCTGAATTCAACTCTGC CATGAACCCCATCATTACTCCTACCGCGACAAAGAAATGAGCGCCACCTTTAGG 10 CAGATCCTCTGCTGCCAGCGCAGTGAGAACCCCACCGGCCCCACAGAAAGCTCA GACCGCTCGGCTTCCCCCCAACCACCATCTTGGCTGGAGTTCACAGCAATG ACCACTCTGTGGTTTAGAACGGAAACTGAGATGAGGAACCAGCCGTCCTCTTG GAGGATAAACAGCCTCCCCTACCCAATTGCCAGGGCAAGGTGGGGTGTGAGAG AGGAGAAAAGTCAACTCATGTACTTAAACACTAACCAATGACAGTATTTGTTCCT 15 GGACCCCACAAGACTTGATATATTGAAAATTAGCTTATGTGACAACCCTCATC TTGATCCCCATCCCTTCTGAAAGTAGGAAGTTGGAGCTCTTGCAATGGAATTCAA GAACAGACTCTGGAGTGTCCATTTAGACTACACTAACTAGACTTTTAAAAGATTT TGTGTGGTTTGGTGCAAGTCAGAATAAATTCTGGCTAGTTGAATCCACAACTTCA TTTATATACAGGCTTCCCTTTTTTATTTTTAAAGGATACGTTTCACTTAATAAACA 20 CGTTTATGCCTATCAGCAAAAAAAAAAAAAAAAA

SEQ ID NO: 697. A STATE OF THE SECOND SECOND

SEQ ID NO: 698

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Human interleukin 11 mRNA, complete cds gi|186272|gb|M57765.1|HUMIL11[186272]
GCTCAGGGCACATGCCTCCCCTCCCCAGGCCGCGCCCAGCTGACCCTCGGGGCT
CCCCCGGCAGCGGACAGGGAAGGGTTAAAGGCCCCCGGCTCCCTGCCCCTGCC
CTGGGGAACCCCTGGCCCTGTGGGGACATGAACTGTGTTTGCCGCCTGGTCCTGG
TCGTGCTGAGCCTGTGGCCAGATACAGCTGTCGCCCCTGGGCCACCACCTGGCCC
CCCTCGAGTTTCCCCAGACCCTCGGGCCGAGCTGGACAGCACCGTGCTCCTGACC
CGCTCTCTCTGGCGGACACGCGGCAGCTGCACAGCTGAGGGACAAATTC
CCAGCTGACGGGGACCACAACCTGGATTCCCTGCCCACCCTGGCCATGAGTGCG
GGGGCACTGGGAGCTCTACAGCTCCCAGGTGTGCTGACAAGGCTGCGAGCGGAC
CTACTGTCCTACCTGCGCACGTGCAGTGCTGCCCGGGCAGTGGCTCCC
TGAAGACCCTGGAGCCCGAGCTGGCCCTGCCCCAGCCACCCCGGA
CCCGCCGGCTGCAGCTCCTGATGTCCCGCCTGGCCCCAGCCACCCCCGGA
CCCGCCGGCGCCCCCCCCTCCTCAGCCTGGGGGGGCATCAGGGCC

GCCACGCCATCCTGGGGGGGGCTGCACCTGACACTTGACTGGGCCGTGAGGGGA CTGCTGCTGCTGAAGACTCGGCTGTGACCCGGGGCCCAAAGCCACCACCGTCCTT CCAAAGCCAGATCTTATTTATTTATTTATTTCAGTACTGGGGGCGAAACAGCCAG GTGATCCCCCGCCATTATCTCCCCCTAGTTAGAGACAGTCCTTCCGTGAGGCCT GGGGGACATCTGTGCCTTATTTATATTTATTTCAGGAGCAGGGGTGGGAGG CAGGTGGACTCCTGGGTCCCCGAGGAGGAGGGGACTGGGGTCCCGGATTCTTGG GTCTCCAAGAAGTCTGTCCACAGACTTCTGCCCTGGCTCTTCCCCATCTAGGCCTG GGCAGGAACATATATTATTTATTTAAGCAATTACTTTTCATGTTGGGGTGGGAC GGAGGGAAAGGGAAGCCTGGGTTTTTGTACAAAAATGTGAGAAACCTTTGTGA

SEQ ID NO: 699

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Homo sapiens mRNA for GABA-BR1a (hGB1a) receptor gil2826760|emblY11044.1|HSGTHLA1[2826760]

gi|2826760|emb|Y11044.1|HSGTHLA1[2826760] 15 CGCAGACCCCCAACGCCACCTCAGAAGGTTGCCAGATCATACACCCGCCCTGGG AAGGGGGCATCAGGTACCGGGGCCTGACTCGGGACCAGGTGAAGGCTATCAACT TCCTGCCAGTGGACTATGAGATTGAGTATGTGTGCCGGGGGGAGCGCGAGGTGG TGGGGCCCAAGTCCGCAAGTGCCTGGCCAACGGCTCCTGGACAGATATGGACA CACCCAGCCGCTGTGTCCGAATCTGCTCCAAGTCTTATTTGACCCTGGAAAATGG 20 GAAGGTTTTCCTGACGGGTGGGGACCTCCCAGCTCTGGACGGAGCCCGGGTGGA TTTCCGGTGTGACCCCGACTTCCATCTGGTGGGCAGCTCCCGGAGCATCTGTAGT CAGGGCCAGTGGAGCACCCCGAAGCCCCACTGCCAGGTGAATCGAACGCCACAC ##CCAGGGGGCCAGGCCTGCCAGCCGGTGGAGATGGCGCTGGAGGACGTGAAT AGCCGCAGGGACATCCTGCCGGACTATGAGCTCAAGCTCATCCACCACGACAGC AAGTGTGATCCAGGCCAAGCCACCAAGTACCTATATGAGCTGCTCTACAACGAC CCTATCAAGATCATCCTTATGCCTGGCTGCAGCTCTGTCTCCACGCTGGTGGCTG AGGCTGCTAGGATGTGGAACCTCATTGTGCTTTCCTATGGCTCCAGCTCACCAGC 30 CCTGTCAAACCGGCAGCGTTTCCCCACTTTCTTCCGAACGCACCCATCAGCCACA

CTCACAACCGGCAGCGTTTCCCCACTTTCTTCCGAACGCACCCATCAGCCACA
CTCCACAACCCTACCCGCGTGAAACTCTTTGAAAAAGTGGGGCTGGAAGAAGATT
GCTACCATCCAGCAGACCACTGAGGTCTTCACTTCGACTCTGGACGACCTGGAGG
AACGAGTGAAGGAGCTGGAATTGAGATTACTTTCCGCCAGAGTTTCTTCTCAGA
TCCAGCTGTGCCCGTCAAAAACCTGAAGCGCCAGGATGCCCGAATCATCGTGGG
35 ACTTTTCTATGAGACTGAAGCCCGGAAAGTTTTTTTGTGAGGTGTACAAGGAGCGT
CTCTTTGGGAAGAAGTACGTCTGGTTCCTCATTGGGTGGTATGCTGACAATTGGT

CTCTTTGGGAAGAAGTACGTCTGGTTCCTCATTGGGTGGTATGCTGACAATTGGT TCAAGATCTACGACCCTTCTATCAACTGCACAGTGGATGAGATGACTGAGGCGGT GGAGGGCCACATCACAACTGAGATTGTCATGCTGAATCCTGCCAATACCCGCAG CATTTCCAACATGACATCCCAGGAATTTGTGGAGAAACTAACCAAGCGACTGAA AAGACACCCTGAGGAGACAGGAGGCTTCCAGGAGGCACCGCTGGCCTATGATGC

40 AAGACACCCTGAGGAGACAGGAGGCTTCCAGGAGGCACCGCTGGCCTATGATGC
CATCTGGGCCTTGGCACTGGCCCTGAACAAGACATCTGGAGGAGGCGGCCGTTCT
GGTGTGCGCCTGGAGGACTTCAACTACAACAACCAGACCATTACCGACCAAATC
TACCGGGCAATGAACTCTTCGTCCTTTGAGGGTGTCTCTGGCCATGTGGTGTTTTG
ATGCCAGCGGCTCTCGGATGGCATGACCAAGGATGATCTTTCCTCCTCCAA

45 GCTACAAGAAGATTGGCTACTATGACAGCACCAAGGATGATCTTTCCTGGTCCAA
AACAGATAAATGGATTGGAGGGTCCCCCCCAGCTGACCAGACCCTGGTCATCAA
GACATTCCGCTTCCTGTCACAGAAACTCTTTATCTCCGTCTCAGTTCTCCAGCC
TGGGCATTGTCCTAGCTGTTGTCTGTCCTTTAACATCTACAACTCACATGTC
CGTTATATCCAGAACTCACAGCCCAACCTGAACAACCTGACTGCTGTGGGCTGCT

CACTGGCTTTAGCTGCTGTCTTCCCCCTGGGGCTCGATGGTTACCACATTGGGAG GAACCAGTTTCCTTTCGTCTGCCAGGCNCGCCTCTGGCTCCTGGGCCTGGGCTTTA GTCTGGGCTACGGTTCCATGTTCACCAAGATTTGGTGGGTCCACACGGGCTTCAC AAAGAAGGAAGAAAGAAGGAGTGGAGGAAGACTCTGGAACCCTGGAAGCTGT 5 ATGCCACAGTGGGCCTGCTGGTGGGCATGTCCTCACTCTCGCCATCTGGCA GATCGTGGACCCTCTGCACCGGACCATTGAGACATTTGCCAAGGAGGAACCTAA GGAAGATATTGACGTCTCTATTCTGCCCCAGCTGGAGCATTGCAGCTCCAGGAAG ATGAATACATGGCTTGGCATTTTCTATGGTTACAAGGGGCTGCTGCTGCTGCTGG GAATCTTCCTTGCTTATGAGACCAAGAGTGTCCCACTGAGAAGATCAATGATCA 10 CCGGGCTGTGGCCATGCTATCTACAATGTGGCAGTCCTGTGCCTCATCACTGCT CCTGTCACCATGATTCTGTCCAGCCAGCAGGATGCAGCCTTTGCCTTTGTCTCT TGCCATAGTTTTCTCCTCCTATATCACTCTTGTTGTGCTCTTTTGTGCCCAAGATGC GCAGGCTGATCACCCGAGGGGAATGGCAGTCGGAGGCGCAGGACACCATGAAG ACAGGGTCATCGACCAACAACAACGAGGAGGAGAAGTCCCGGCTGTTGGAGAA 15 GGAGAACCGTGAACTGGAAAAGATCATTGCTGAGAAAGAGGAGCGTGTCTCTGA ACACCCCAGAACCCTCTGGGGGCCTGCCCAGGGGACCCCCTGAGCCCCCGAC CGGCTTAGCTGTGATGGGAGTCGAGTGCATTTGCTTTATAAGTGAGGGTAGGGTG AGGGAGGACAGGCCAGTAGGGGGAGGGGAAAGGGAGAGGGGAAGGGCAGGGGA 20 CTCAGGAAGCAGGGGTCCCCATCCCAGCTGGGAAGAACATGCTATCCAATCT CATCTCTTGTAAATACATGTCCCCCTGTGAGTTCTGGGCTGATTTGGGTCTCTCAT ACGGCAACCCCTGCAGCTCCTCTGCCTTTGTGCTCTGTTCCTGTCCAGCAGGGGTC ::25 TCCCAACAAGTGCTCTTTCCACCCCAAAGGGGCCTCTCCTTTTCTCCACTGTCATA ATCTCTTTCCATCTTACTTGCCCTTCTATACTTTCTCACATGTGGCTCCCCCTGAAT TTTGCTTCCTTTGGGAGCTCATTCTTTTCGCCAAGGCTCACATGCTCCTTGCCTCT GCTCTGTGCACTCACGCTCAGCACACATGCATCCTCCCCTCTCCTGCGTGTGCCCA 30 CTGAACATGCTCATGTTACACACGCTTTTCCCGTATGCTTTCTTCATGTTCAGTC ACATGTGCTCTCGGGTGCCCTGCATTCACAGCTACGTGTGCCCCTCTCATGGTCAT GGGTCTGCCCTTGAGCGTGTTTGGGTAGGCATGTGCAATTTGTCTAGCATGCTGA GTCATGTCTTTCCTATTTGCACACGTCCATGTTTATCCATGTACTTTCCCTGTGTAC CCTCCATGTACCTTGTGTACTTTCTTCCCTTAAATCATGGTATTCTTCTGACAGAG 35 CCATATGTACCCTACCCTGCACATTGTTATGCACTTTTCCCCAATTCATGTTTGGT GGGGCCATCCACACCCTCTCCTTGTCACAGAATCTCCATTTCTGCTCAGATTCCCC CCATCTCCATTGCATTCATGTACTACCCTCAGTCTACACTCACAATCATCTTCTCC CAAGACTGCTCCCTTTTGTTTTTGTGTTTTTTTGAGGGGAAATTAAGGAAAAATAAG TGGGGCAGGTTTGGAGAGCTGCTTCCAGTGGATAGTTGATGAGAATCCTGACC 40 AAAGGAAGCACCCTTGACTGTTGGGATAGACAGATGGACCTATGGGGTGGGAG GTGGTGTCCCTTTCACACTGTGGTGTCTCTTGGGGAAGGATCTCCCCGAATCTCA

SEQ ID NO: 700

zh96g08.s1 Soares_fetal_liver_spleen_1NFLS_S1 Homo sapiens cDNA clone IMAGE:429182 3', mRNA sequence gi|1448327|gb|AA004759.1|AA004759[1448327] ACTTTATGCAAAAAAAAAAATATACATTTATTTATAGGTCTCAATACAGCAAAATGA AAACGAAAATTGAGAACATTGCTCATTAGGCCAGCAACTTTAAAATTATTTAATT TGAAATATAAAATAGGTGGTCTTCATAAAAAAGATGCATGAAATTACCTTACCTT

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SEQ ID NO: 701

Homo sapiens canalicular multispecific organic anion transporter 2 (CMOAT2) mRNA, complete cds gi|3550323|gb|AF083552.1|AF083552[3550323]

- 25 GGGTGCCGGCCCAGGCCCCGGAAAAAATGCCTCCGGCGAGGACGAGGTGCTGCTC
 TTCGGCTCCAGCTCCTCATCAGTGCCTGCTTCAAGCTTATCCAGGACCTGCTCTC
 CTTCATCAATCCACAGCTGCTCAGCATCCTGATCAGGTTTATCTCCAACCCCATG
 GGCCCCTCCTGGTGGGGCTTCCTGGTGGCTGATCATCTTCTGTGCTCCATGA
 TGCAGTCGCTGATCTTACAACACTATTACCACTACATCTTTGTGACTGGGTGAA
 30 GTTTCGTACTGGGATCATGGGTGTCATCTACAGGAAGGCTCTGGTTATCACCAAC
 - TCAGTCAAACGTGCGTCCACTGTGGGGGAAATTGTCAACCTCATGTCAGTGGATG
 CCCAGCGCTTCATGGACCTTGCCCCCTTCCTCAATCTGCTGTGGTCAGCACCCCTG
 CAGATCATCCTGGCGATCTACTTCCTCTGGCAGAACCTAGGTCCCTCTGTCCTGG
 CTGGAGTCGCTTTCATGGTCTTGCTGATTCCACTCAACGGAGCTGTGGCCGTGAA
 35 GATGCGCGCCTTCCAGGTAAAGCAAATGAAATTGAAGGACTCGCGCATCAAGCT
 GATGAGTGAGATCCTGAACGGCATCAAGGTGCTGAAGCTGTACGCCTGGGAGCC

 - 40 TGGACGCCGAGAAGGCCTTTGTGTCTGTGTCCTTGTTTAATATCTTAAGACTTCCC CTCAACATGCTGCCCCAGTTAATCAGCAACCTGACTCAGGCCAGTGTGTCTCTGA AACGGATCCAGCAATTCCTGAGCCAAGAGGAACTTGACCCCCAGAGTGTGGAAA GAAAGACCATCTCCCCAGGCTATGCCATCACCATACACAGTGGCACCTTCACCTG GGCCCAGGACCTGCCCCCCACTCTGCACAGCCTAGACATCCAGGTCCCGAAAGG

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GAGAGAAGGCATTAACCTGTCTGGGGGCCAGCGGCAGCGGGTCAGTCTGGCTC GAGCTGTTTACAGTGATGCCGATATTTTCTTGCTGGATGACCCACTGTCCGCGGT GGACTCTCATGTGGCCAAGCACATCTTTGACCACGTCATCGGGCCAGAAGGCGTG CTGGCAGGCAAGACGCGAGTGCTGGTGACGCACGGCATTAGCTTCCTGCCCCAG ACAGACTTCATCATTGTGCTAGCTGATGGACAGGTGTCTGAGATGGGCCCGTACC CAGCCTGCTGCAGCGCAACGGCTCCTTTGCCAACTTTCTCTGCAACTATGCCCC CGATGAGGACCAAGGGCACCTGGAGGACAGCTGGACCGCGTTGGAAGGTGCAG AGGATAAGGAGCACTGCTGATTGAAGACACACTCAGCAACCACACGGATCTGA CAGACAATGATCCAGTCACCTATGTGGTCCAGAAGCAGTTTATGAGACAGCTGA 10 GTGCCTGTCCTCAGATGGGGAGGGACAGGGTCGGCCTGTACCCCGGAGGCACC TGGGTCCATCAGAGAAGGTGCAGGTGACAGAGGCGAAGGCAGATGGGGCACTG ACCCAGGAGGAGAAAGCAGCCATTGGCACTGTGGAGCTCAGTGTGTTCTGGGAT TATGCCAAGGCCGTGGGGCTCTGTACCACGCTGGCCATCTGTCTCCTGTATGTGG GTCAAAGTGCGGCTGCCATTGGAGCCAATGTGTGGCTCAGTGCCTGGACAAATG ATGCCATGGCAGACAGTAGACAGAACACTTCCCTGAGGCTGGGCGTCTATG CTGCTTTAGGAATTCTGCAAGGGTTCTTGGTGATGCTGGCAGCCATGGCCATGGC AGCGGGTGGCATCCAGGCTGCCCGTGTGTTGCACCAGGCACTGCTGCACAACAA GATACGCTCGCCACAGTCCTTCTTTGACACCACCACCATCAGGCCGCATCCTGAAC TGCTTCTCCAAGGACATCTATGTCGTTGATGAGGTTCTGGCCCCTGTCATCCTCAT 20 GCTGCTCAATTCCTTCTTCAACGCCATCTCCACTCTTGTGGTCATCATGGCCAGCA CGCCGCTCTTCACTGTGGTCATCCTGCCCCTGGCTGTGCTCTACACCTTAGTGCAG AND CONCACCIATCIACTCCCACTITICGGAGACAGIGACIGGIGCCAGIGICATCCGGGC ********CTACAACCGCAGCCGGATTTTGAGATCATCAGTGATACTAAGGTGGATGCCAA CCAGAGAAGCTGCTACCCCTACATCTCCCAACCGGTGGCTGAGCATCGGAGTG GAGTTCGTGGGAACTGCGTGGTGCTCTTTGCTGCACTATTTGCCGTCATCGGGA GGAGCAGCCTGAACCCGGGGCTGGTGGGCCTTTCTGTGTCCTACTCCTTGCAGGT GACATTTGCTCTGAACTGGATGATACGAATGATGTCAGATTTGGAATCTAACATC GTGGCTGTGGAGAGGGTCAAGGAGTACTCCAAGACAGAGACAGAGGCGCCCTGG 30 GTGGTGGAAGGCAGCCCCCCCGAAGGTTGGCCCCCACGTGGGGAGGTGGAG TTCCGGAATTATTCTGTGCGCTACCGGCCGGGCCTAGACCTGGTGCTGAGAGACC TGAGTCTGCATGTGCACGGTGGCGAGAAGGTGGGGATCGTGGGCCGCACTGGGG CTGGCAAGTCTTCCATGACCCTTTGCCTGTTCCGCATCCTGGAGGCGGCAAAGGG TGAAATCCGCATTGATGGCCTCAATGTGGCAGACATCGGCCTCCATGACGTGCGC TCTCAGCTGACCATCATCCCGCAGGACCCCATCCTGTTCTCGGGGACCCTGCGCA TGAACCTGGACCCCTTCGGCAGCTACTCAGAGGAGGACATTTGGTGGGCTTTGGA GCTGTCCCACCTGCACACGTTTGTGAGCTCCCAGCCGGCAGGCCTGGACTTCCAG TGCTCAGAGGGCGGGAGAATCTCAGCGTGGGCCAGAGGCAGCTCGTGTGCCTG GCCCGAGCCTGCTCCGCAAGAGCCGCATCCTGGTTTTAGACGAGGCCACAGCTG 40 CCATCGACCTGGAGACTGACAACCTCATCCAGGCTACCATCCGCACCCAGTTTGA TACCTGCACTGTCCTGACCATCGCACACCGGCTTAACACTATCATGGACTACACC AGGGTCCTGGTCCTGGACAAAGGAGTAGTAGCTGAATTTGATTCTCCAGCCAACC TCATTGCAGCTAGAGGCATCTTCTACGGGATGGCCAGAGATGCTGGACTTGCCTA GACACCAAATATGTCCGCAGAATGGACTTGATAGCAAACACTGGGGGCACCTTA AGATTTTGCACCTGTAAAGTGCCTTACAGGGTAACTGTGCTGAATGCTTTAGATG AGGAAATGATCCCCAAGTGGTGAATGACACGCCTAAGGTCACAGCTAGTTTGAG CCAGTTAGACTAGTCCCCGGTCTCCCGATTCCCAACTGAGTGTTATTTGCACACT

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SEQ ID NO: 702

yq42d10.s1 Soares fetal liver spleen 1 NFLS Homo sapiens cDNA clone IMAGE:198451 3', mRNA sequence gi|970054|gb|R94659.1|R94659[970054]

TTGTTTTTTTGGTTCAGCATAACTTGGAACATTTGAAAGCTTTTCAACCTAAATG

10 TGGG

GAAAAAACAGGTAAGGCATTATTTTTGCACAAAACTAGCATTCCTAATAGTGCA AATGAA

 ${\tt TCTGATACCTCTTAAAATGGTGAGAGGTCATACACTTACTAGATTAGATT}\\ {\tt TTCTT}$

15 TCTATGGCTTGACAAATTATCCCTCTATAAATTCTACTCTCACCCAGAGGCTGTTG CTGT

AATCAAAAGGATAACTGTAGGATAAAGGTCCAACCTTCTCCTGGTATCCGGCAA AAGGGT

TTTTGCTCATATGGCAAAAAAAATCTAATTTTTAAATTATCCTACAGNGGAATAT

20 ACAAC

TGGGNTTCCTNGGGACCCTCTATTTATCNGGCGGCAACAGGTGGTTCGGGGCGGC

TO WISEQ ID NO: 703 White the control of the contro

25 zd29f03.s1 Soares_fetal_heart_NbHH19W Homo sapiens cDNA clone IMAGE:342077 3', mRNA sequence gi|1367074|gb|W60315.1|W60315[1367074] CATAACTTAAGTAAACTTTATTTTCAAAAATGCTTCAGGTACAAAAGAAAAACAATC GGCAAAGTCTAACAATAATTAACAAACCAGCTCTTGAGCGGCAGAGTGCTCCAG GGATGAGAGGGGCTGGGGATGGAAAGGTGGTTGGGAGACACAACATTTTTCTAG

DESCRIPTION REPORTS THE LIGHT ARM IN THE FRANCE STREET WAS LESS BUT IN THE PROPERTY OF THE PRO

30 CTTCAGAAAGTCAGGGAGCCCAGATCACAGCCTGAACTTCATGGTATTGGTTACA GATTCTTTACAAAGGTGTTTACCTCTCTCATGAGGTCTTCTTGATTGGTTACTTCC TCAGAAAAATCATCATTGACATCCAACACCAGCACTGGAATGTTCATCAGAGCCT CAAAGTGGAGCCTGTCACTTGTACACANGACCTCTCAAAGATCTGTACTGGCTTC CTGGCCTGGTAAGAGTTCTCAGGGGAAG

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SEQ ID NO: 704

yb54f05.r1 Stratagene ovary (#937217) Homo sapiens cDNA clone IMAGE:75009 5', mRNA sequence gi|653755|gb|T51895.1|T51895[653755]

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SEQ ID NO: 705

zx69a01.s1 Soares_total_fetus_Nb2HF8_9w Homo sapiens cDNA clone IMAGE:796680 3', mRNA sequence gi|2185799|gb|AA460679.1|AA460679[2185799] TACTCAGTCACCACCCAGAAATTGTCCGAGTTATGAAAATAGATTCATTTTGAGAA

SEQ ID NO: 706

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- zv64g11.s1 Soares_total_fetus_Nb2HF8_9w Homo sapiens cDNA clone IMAGE:758468 3', mRNA sequence gi|2046825|gb|AA393856.1|AA393856[2046825]
 TTTAACATCAGTTAAAGATTTTATTTGATTCATTAAAGAGGAAACTGGTGAGGCA TTTCCACCAGCTCAAGGAAGAATTTTGTAAATGTTATATTTATGGATCAGAAATA ACTGAAATGAATGTGCAAATGGAGGCAAAACTGGCCTCTTCCACAGTGGGGAAG
 AAAGTCAACAGAACCTCCACTAGGCATAATTTACATATGTACAGACTCAATCAGC
- 15 AAAGTCAACAGAACCTCCACTAGGCATAATTTACATATGTACAGACTCAATCAGC
 TTTTAATATAGAAAGATATTTGAACCCAAAATCTTTCATTAAGGTAAAAAATACA
 ATAATAATTTTTAATGAAATCCTGGAAAATTCATACAAATAAAATTAAAAGCCTC
 CAATGGGGTATAATCCAGCAATATCCTAGGCAAATGCCTCCTGAAGAACAACAG
 CCTTTTTAAAAACATCACTGTTTATCATTCAAAAATTCAGACGTCTCCTATCTTTGGC
- 20 TATTTTATCTCTTCAACT

aa47b01.r1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:824041 5! similar to TR:G1049078 G1049078 SRP30C:;, mRNA sequence

- 25 gi|2219894|gb|AA490721.1|ÆA490721[2219894]
 TATCTCAGAAAAGACATGCGATATGCCCTGCGTAAACTGGATGACACCAAA
 TTCCGCTCTCATGAGGGTGAAACTTCCTACATCCGAGTTTATCCTGAGAGAAGCA
 CCAGCTATGGCTACTCACGGTCTCGGTCTAGGTCAAGGGGCCGTGACTCTCCATA
 CCAAAGCAGGGGTTCCCACACTACTTCTCTCTCTTCAGGCCCTACTGAGACAGGT

SEQ ID NO: 708

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ACCTACTCCTGGTAAGTGGGGTTGCGGATGAGGGGGACGGGGCGTGGCGCTGGC TGGCGTGAGAAGTGCGGTGCTGATGTCCCTCTGTCGGGTTTTTGCAGCGTCGGCG TGTTCAAGAACGCCGCGTGGAGATCATCGCCAACGATCAGGGCAACCGCATCA CGCCGTCCTATGTCGCCTTCACTCCTGAAGGGGAACGTCTGATTGGCGATGCCGC CAAGAACCAGCTCACCTCCAACCCGAGAACACGGTCTTTGACGCCAAGCGGCT CATCGGCCGCACGTGGAATGACCCGTCTGTGCAGCAGGACATCAAGTTCTTGCCG TATTTAGAGTTATAAGTCTCTGGAAAAGTGTTGAGACAACAGTTGAAGGTTATAG ACATGATGTATGTAATAACTTTAATACTATTAGTATGTTACAAAACTTAAGACAG 10 TTGCTGTCGTACTGTCTACGATAGTTTAGGAATAAAAGACCGATTAAAACTGAAC TTTGTAAGACACCTATACTCCCTGAAGTATTTCTAGTCAATTTGCAGCCCCAAGG GACCAAAATAAACCAAATTGTGGGGATGGTAGTGGGTCTTTTAAACTTTGAGATG TCATTGTATCTGTGTCTGAAAACAATAATTCTTTAAAATAGGTGGTTGAAAAGAA TATTTGGGAAAGAAGGTAAATATTTCTAGAACAATGTTAAGTATTTTTTGATCAT TAGTATTCTCGGTTGGCTGTTATGTATAGAAGCCTTCGTGAAGGGTTTCAAAAAT TTTAATCAGAATGGTATTCATGCTTGTCACGGTTTAATTATTGAGTCCCTTTACTA TAAGCCAAACAAAATAGACTTTTCATGTATTATTTAATGCTTACAATTCCAGGA 20 ACAATAAAATTTTATATGTTGTATTCATCAATAATTGGCTTAAAAACTAAAGTGA TGGTTTGACTGTAATTTTTTTTTTTGAGATGGAGTCTTGCTCTGTTGCCCAGGCT GGACTGCAGTGCACGATCTCAGCTCACTGCAACCTCTGCCTCCCGGGTTAAGCA SOME SECURITION OF THE PROPERTY OF THE PROPERT GCTAATTTTTTTTTTTTTTTAATTTTCAGTAGAGAGAGGGTTTCTCCACATTGCC AGGCTGGTCTTGAAATCCTGCCCTCAGGTTGATCCTCCTGCCTAGCCTCCCAAAG TGCTGGATTATAGGCAGAAGCCACCGCCTGGCCAGACTGTAATTTAAATAAGGG TTAAACTATGTGACAATACACTTAATTATCTTTATCCTTTTAGGTTACCCATGCAG TTGTTACTGTACCAGCCTATTTTAATGATGCCCAACGCCAAGCAACCAAAGACGC TGGAACTATTGCTGGCCTAAATGTTATGAGGATCATCAACGAGCCGTAAGTATGA 30 AATTCAGGGATACGGCATATTTGCCAAATAGTGGAAATGTGAAGTACTGACAAA ACTTTTCCCTTTTTCAATCTAATAGTACGCCAGCTGCTATTGCTTATGGCCTGGAT AAGAGGGGGGGGAGAAGAACATCCTGGTGTTTGACCTGGGTGGCGGAACCTTC GATGTGTCTCTCACCATTGACAATGGTGTCTTCGAAGTTGTGGCCACTAATG GAGATACTCATCTGGGTGGAGAAGACTTTGACCAGCGTGTCATGGAACACTTCAT 35 TGCAGAAACTCCGGCGCGAGGTAGAAAAGGCCAAGGCCCTGTCTTCTCAGCATC AAGCAAGAATTGAAATTGAGTCCTTCTATGAAGGAGAAGACTTTTCTGAGACCCT GACTCGGGCCAAATTTGAAGAGCTCAACATGGTATGTTCCTTGTTTTCTGCTTTGC TAATGAGATCTCCTTAGACTCTGAATTCAGGACATTGCATCTAGATACTTAGATA 40 ACAGACATCACAGTAACCATGTCTTTTTTCTAGGATCTGTTCCGGTCTACTATGAA GCCCGTCCAGAAAGTGTTGGAAGATTCTGATTTGAAGAAGTCTGATATTGATGAA ATTGTTCTTGTTGGTGGCTCGACTCGAATTCCAAAGATTCAGCAACTGGTTAAAG AGTTCTTCAATGGCAAGGAACCATCCCGTGGCATAAACCCAGATGAAGCTGTAG CGTATGGTGCTGTCCAGGCTGGTGTGCTCTCTGGTGATCAAGATACAGGTAG GTCATCATCGCAGCATCTTTCTTAGTGATTCAGTAGCTTGATGGAAGAGCTCGGT ACCCCTATTGCTTTAGAAAATACCAGAATATGAGCAACAAGGTCACACAGCTAG TAAAGGGTATAAGTGAAGACAAGACTGGGGTAGTCTCCAAGATCATTAGCAACT GTTTAATTCACTGCCTTTAAAATGTGTGTGTTAGAACCTAACCAAATGTTAGAGA GATAAACTTTACATAGCTCATAGGGAGAACTTGAATTAAAAGTTAAATAACTTAT

CCTTACAGGTGACCTGGTACTGCTTCATGTATGTCCCCTTACACTTGGTATTGAAA CTGTAGGAGGTGTCATGACCAAACTGATTCCAAGTAATACAGTGGTGCCTACCAA 5 GAAGTCTTGCTCTGTTGCCCAGGCTGGACTGCAGTGGCACGATCTCGGCTCACTG CAAATTCTGTCTCCCGGGTTCAAGTGATTCTCCTGCCTCAGCCTCCAGAGTAGCT GGATTACAGCCTGACCACCACACCTGGCTAATTTCTGTATTTTTAGTAGAGGATG GGCTTTCACCATGTTTCCCAGGCTGGTCTCCAACTCCTGACCTCAGGTCATCTGCC TGCCTCCACCGTCCCGAAAGTACTGGGATTATAGCGTGAGCCACCACGCCAGATC 10 TATCTATCATGGCATATTTTAAAAGAACATGACTTAATATGTCCTATTGAAATGG CTAGGGAACTAAGTAACTGCTGTTTTCAGATGGAGGTCTTAATTTGAATAATGTT GATATTAGATATTTAGCATTCTTTTTTTTTTTTTTAATGGAGTCTTGCTCTGTCG CCTAGGCTGGGTGCAGTGGCATGACTTGCAACCTCTGCCTCCCGAATAGCTGGG ATTACAGGTGCCCACCATCACGCCCGGCTAAGTTTTGTATTTTTAGTAGAGGCGA 15 GTTTCGCCATGTTGGCCAGGCTGGTCTTGAACCCCTAACCTCAGTGATCCCACGG TCACCGACCTGGCCTCCCAAAAGTACTGTACCCAGCCAATGATTAGCATTCTCAC TAATAATAGCATCTGAGCTGGCTCCTAGAGTACAAGAAAAAGGAGTTCACAGTA CTTTAAAATAGATAAAATTCAGTTGAGTTAGTAACCTAACTCATTGTTAGTACTA GTTGCTGCTCCTTGTAGACCAATATGAAATTACTTTTAGCTCGATAAAACCAAAA 20 GTGTCACTTTATGCTTCAGACTGAAATGCGGGGATCTAGATGTGCTAATGCTTGT CAGTAACAACTAACAAGTTTTTCTGTATGTAACTTCTAGGTGAAAGACCCCTGAC TAYAAAGACAATCATCTTCTGGGTACATTTGATCTGACTGGAATTCCTCCTGCTCCTC SECOND CONTROL OF THE AGTGACAGCTGAAGACAAGGGTACAGGGXACAAAAATAAGATCACAATCACCA 25 ATGACCAGAATCGCCTGACACCTGAAGAAATCGAAAGGATGGTTAATGATGCTG AGAAGTTTGCTGAGGAAGACAAAAAGCTGAAGGAGCGCATTGATACTAGAAATG AGTTGGAAAGCTATGCCTATTCTCTAAAGAATCAGATTGGAGATAAAGAAAAGC TGGGAGGTAAACTTTCCTCTGAAGATAAGGAGACCATGGAAAAAGCTGTAGAAG AAAAGATTGAATGGCTGGAAAGCCACCAAGATGCTGACATTGAAGACTTCAAAG 30 CTAAGAAGAAGGAACTGGAAGAAATTGTTCAACCAATTATCAGCAAACTCTATG GAAGTGCAGGCCCTCCCCCAACTGGTGAAGAGGGATACAGCAGAAAAAGATGAGT TGTAGACACTGATCTGCTAGTGCTGTAATATTGTAAATACTGGACTCAGGAACTT TTGTTAGGAAAAATTGAAAGAACTTAAGTCTCGAATGTAATTGGAATCTTCACC TCAGAGTGGAGTTGAAACTGCTATAGCCTAAGCGGCTGTTTACTGCTTTTCATTA 35 GCAGTTGCTCACATGTCTTTGGGTGGGGGGGAGAAGAAGAATTGGCCATCTTAA AAAGCGGGTAAAAACCTGGGTTAGGGTGTGTGTCACCTTCAAAATGTTCTATT TAACAACTGGGTCATGTGCATCTGGTGTAGGAGGTTTTTTCTACCATAAGTGACA CCAATAAATGTTTGTTATTTACACTGGTCTAATGTTTGTGAGAAGCTT

40 SEO ID NO: 709

Human adenosine receptor (A2) gene, complete cds
gi|177891|gb|M97370.1|HUMA2XXX[177891]
GGCACGAGGCTGGCTGAGCCATGATGCTGCCAGAACCCCTGCAGAGGGCCT
GGTTTCAGGAGACTCAGAGTCCTCTGTGAAAAAAGCCCTTGGAGAGGCGCCCCAG
45 CAGGGCTGCACTTGGCTCCTGTGAGGAAGGGGCTCAGGGTCTGGGCCCCTCCGCC
TGGGCCGGGCTGGGAGCCAGGCGGGCGGCTGCAGCAATGGACCGTGAGC
TGGCCCAGCCCGCGTCCGTGCTGAGCCTGCCTGTCGTCTGTGCCATCAT
GGGCTCCTCGGTGTACATCACGGTGGAGCTGGCCATTGCTGTGCCATCCTG
GGCAATGTGCTGGTGTGCTGGGCCGTTCTGTGCCAGAACGTC

ACCAACTACTTGTGGTGTCACTGGCGGCGGCCGACATCGCAGTGGGTGTGCTCG CCTCTTCATTGCCTGCTTCGTCCTGGTCCTCACGCAGAGCTCCATCTTCAGTCTCC TGGCCATCGCCATTGACCGCTACATTGCCATCCGCATCCCGCTCCGGTACAATGG 5 CTTGGTGACCGGCACGAGGGCTAAGGGCATCATTGCCATCTGCTGGGTGCTGTCG TTTGCCATCGCCTGACTCCCATGCTAGGTTGGAACAACTGCGGTCAGCCAAAGG AGGGCAAGAACCACTCCCAGGGCTGCGGGGAGGGCCAAGTGGCCTGTCTCTTTG AGGATGTGGTCCCCATGAACTACATGGTGTACTTCAACTTCTTTGCCTGTGTGCTG GTGCCCTGCTGCTCATGCTGGTGTCTATTTGCGGATCTTCCTGGCGCGCGAC 10 GACAGCTGAAGCAGATGGAGAGCCAGCCTCTGCCGGGGGAGCGGGCACGGTCCA CACTGCAGAAGGAGGTCCATGCTGCCAAGTCACTGGCCATCATTGTGGGGCTCTT TGCCCTCTGCTGGCTGCCCCTACACATCATCAACTGCTTCACTTTCTTCTGCCCCG ACTGCAGCCACGCCCTCTCTGGCTCATGTACCTGGCCATCGTCCTCTCCCACACC AATTCGGTTGTGAATCCCTTCATCTACGCCTACCGTATCCGCGAGTTCCGCCAGA 15 CCTTCCGCAAGATCATTCGCAGCCACGTCCTGAGGCAGCAAGAACCTTTCAAGGC AGCTGGCACCAGTGCCCGGGTCTTGGCAGCTCATGGCAGTGACGGAGAGCAGGT CAGCCTCCGTCTCAACGCCACCCGCCAGGAGTGTGGGCCAACGGCAGTGCTCC TGCCCAAGAGTCCCAGGGGAACACGGGCCTCCCAGACGTGGAGCTCCTTAGCCA 20 TGAGCTCAAGGGAGTGTGCCCAGAGCCCCCTGGCCTAGATGACCCCCTGGCCCA GGATGGAGCAGGAGTGTCCTGATGATTCATGGAGTTTGCCCCTTCCTAAGGGAAG AND A TIGGORAGGACCCTGAGGGCAGCCGGTTCCTACTTTGGACTGAGAGAAGGGAGC Manageceaggctggagcageatgaggeecagcaagaaggcttgggttetgaggaage 25 'AGATGTTTCATGCTGTGAGGCCTTGCACCAGGTGGGGGCCACAGCACCAGC'AGC ATCTTTGCTGGGCAGGGCCCAGCCCTCCACTGCAGAAGCATCTGGAAGCACCACC TTGTCTCCACAGAGCAGCTTGGGCACAGCAGACTGGCCTGAGACTGGG GAGTGGCTCCAACAGCCTCCTGCCACCCACACCACTCTCCCTAGACTCTCCTA GGGTTCAGGAGCTGCTGGGCCCAGAGGTGACATTTGACTTTTTCCAGGAAAAAT GTAAGTGTGAGGAAACCCTTTTATTTATTACCTTTCACTCTCTGGCTGCTGGGT 30 CTGCCGTCGGTCCTGCTAACCTGGCACCAGAGCCTCTGCCGGGGAGCCTCAG GCAGTCCTCTCCTGCTGTCACAGCTGCCATCCACTTCTCAGTCCCAGGGCCATCTC TTGGAGTGACAAAGCTGGGATCAAGGACAGGAGTTGTAACAGAGCAGTGCCAG AGCATGGGCCCAGGTCCCAGGGGAGAGGTTGGGGCTGGCAGGCCACTGGCATGT 35 GCTGAGTAGCGCAGAGCTACCCAGTGAGAGGCCTTGTCTAACTGCCTTTCCTTCT AAAGGGAATGTTTTTTCTGAGATAAAATAAAACGAGCCACATCGTGTTTTAAG CTTGTCCAAATGAAAAAAAAAAAAAAAAAAA

SEQ ID NO: 710

NAAGCCTGGTAAGAATTGGGGGGAACCCACTTGGTATTGNCCCTCTTCCAGGATT TTGGAAATTCCAACCGGCCTTGGNTTTAAGAGAAAANAAGGGNTGGTTCCCACT AAT

5 SEQ ID NO: 711

ab36c08.r1 Stratagene HeLa cell s3 937216 Homo sapiens cDNA clone IMAGE:842894 5' similar to TR:G1256802 G1256802 SODIUM/POTASSIUM-TRANSPORTING ATPASE BETA-3 SUBUNIT.; mRNA sequence gi|2218877|gb|AA489275.1|AA489275[2218877] CTGGCCGAGTGGAAGCTCTTCATCTACAACCCGACCACCGGAGAATTCCTGGGGC

- 10 GCACCGCAAGAGCTGGGGTTTGATCTTGCTCTTCTACCTAGTTTTTATGGGTTCC
 TGGCTGCACTCTTCTCATTCACGATGTGGGTTATGCTTCAGACTCTCAACGATGA
 GGTTCCAAAAATACCGTGACCAGATTCCTAGCCCAGGACTCATGGTTTTTCCAAAA
 CCAGTGACCGCATTGGAATATACATTCAGTAGGTCTGATCCAACTTCGTATGCAG
 GGTACATTGAAGACCTTAAGAAGTTTCTAAAACCATATACTTTAGAAGAACAGA

SEQ ID NO: 712

- 20 za24e08.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:293510 3', mRNA sequence gi|1225735|gb|N69574.1|N69574[1225735]
- 25 TGGCAGTGAATAGAACAGTGATTGTTCATACTACTTGGATCTACTGCCTTAATTT ATACTAGGATGTCAATCCACCATTGATTTTGGACCATCAGTGCCAATGTCNACGT AGCCAAAAAGGCCAAT

SEQ ID NO: 713

- Human mRNA for gamma-interferon inducible early response gene (with homology to platelet proteins) gi|33917|emb|X02530.1|HSINFGER[33917]

 GAGACATTCCTCAATTGCTTAGACATATTCTGAGCCTACAGCAGAGGAACCTCCA
 GTCTCAGCACCATGAATCAAACTGCGATTCTGATTTGCTGCCTTATCTTTCTGACT
 CTAAGTGGCATTCAAGGAGTACCTCTCTCTAGAACCGTACGCTGTACCTGCATCA
- 35 GCATTAGTAATCAACCTGTTAATCCAAGGTCTTTAGAAAAACTTGAAATTATTCC TGCAAGCCAATTTTGTCCACGTGTTGAGATCATTGCTACAATGAAAAAGAAGGGT GAGAAGAGATGTCTGAATCCAGAATCGAAGGCCATCAAGAATTTACTGAAAGCA GTTAGCAAGGAAATGTCTAAAAGATCTCCTTAAAACCAGAGGGGAGCAAAATCG ATGCAGTGCTTCCAAGGATGGACCACACAGAGGCTGCCTCTCCCATCACTTCCCT
- 40 ACATGGAGTATATGTCAAGCCATAATTGTTCTTAGTTTGCAGTTACACTAAAAGG
 TGACCAATGATGGTCACCAAATCAGCTGCTACTACTCCTGTAGGAAGGTTAATGT
 TCATCATCCTAAGCTATTCAGTAATAACTCTACCCTGGCACTATAATGTAAGCTCT
 ACTGAGGTGCTATGTTCTTAGTGGATGTTCTGACCCTGCTTCAAATATTTCCCTCA
 CCTTTCCCATCTTCCAAGGGTACTAAGGAATCTTTCTGCTTTGGGGTTTATCAGAA
- 45 TTCTCAGAATCTCAAATAACTAAAAGGTATGCAATCAAATCTGCTTTTTAAAGAA TGCTCTTTACTTCATGGACTTCCACTGCCATCCTCCCAAGGGGCCCAAATTCTTTC AGTGGCTACCTACATACAATTCCAAACACATACAGGAAGGTAGAAATATCTGAA AATGTATGTGTAAGTATTCTTATTTAATGAAAGACTGTACAAAGTATAAGTCTTA GATGTATATATTTCCTATATTGTTTTCAGTGTACATGGAATAACATGTAATTAAGT

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SEQ ID NO: 714

ab21g06.r1 Stratagene lung (#937210) Homo sapiens cDNA clone IMAGE:841498 5' similar to gb:X54304 MYOSIN REGULATORY LIGHT CHAIN 2, NONSARCOMERIC (HUMAN);, mRNA sequence gi|2217534|gb|AA487370.1|AA487370[2217534]

SEQ ID NO: 715

20 H.sapiens mRNA for central cannabinoid receptor gi|736236|emb|X81120.1|HSCANN6[736236]

HORACOPT CONTROL TO THE CONTROL OF T *****CATTTTGAGCTCAGCCTAATCAAAGACTGAGGTTATGAAGTCGATCCTAGATGGC AN CONTROCAGATACEACETTCCGGACCATCACEAGTGACCTCCTGTACGTGGGCTCAA . 25 ATGACATTCAGTACGAAGACATCAAAGGTGACATGGCATCCAAATTAGGGTACT TCCCACAGAAATTCCCTTTAACTTCCTTTAGGGGAAGTCCCTTCCAAGAGAAGAT GACTGCGGGAGACACCCCCAGCTAGTCCCAGCAGACCAGGTGAACATTACAGA ATTTTACAACAAGTCTCTCTCGTCCTTCAAGGAGAATGAGGAGAACATCCAGTGT CTGGCCATTGCAGTCCTGTCCCTCACGCTGGGCACCTTCACGGTCCTGGAGAACC 30 TCCTGGTGCTGTGCGTCATCCTCCACTCCCGCAGCCTCCGCTGCAGGCCTTCCTAC CACTTCATCGGCAGCCTGGCGGTGGCAGACCTCCTGGGGAGTGTCATTTTTGTCT ACAGCTTCATTGACTTCCACGTGTTCCACCGCAAAGATAGCCGCAACGTGTTTCT GTTCAAACTGGGTGGGGTCACGGCCTCCTTCACTGCCTCCGTGGGCAGCCTGTTC 35 CTCACAGCCATCGACAGGTACATATCCATTCACAGGCCCCTGGCCTATAAGAGGA TTGTCACCAGGCCCAAGGCCGTGGTGGCGTTTTGCCTGATGTGGACCATAGCCAT TGTGATCGCCGTGCTCCCTCGGGCTGGAACTGCGAGAAACTGCAATCTGTT TGCTCAGACATTTTCCCACACATTGATGAAACCTACCTGATGTTCTGGATCGGGG

45 GTGAACCCCATCATCTATGCTCTGAGGAGTAAGGACCTGCGACACGCTTTCCGGA GCATGTTTCCCTCTTGTGAAGGCACTGCGCAGCCTCTGGATAACAGCATGGGGGA CTCGGACTGCCTGCACAAACACGCAAACAATGCAGCCAGTGTTCACAGGGCCGC AGAAAGCTGCATCAAGAGCACAGTCAAGATTGCCAAGGTAACCATGTCTGTGTC CACAGACACGTCTGCCGAGGCTCTGTGAGCCTGATGCCTCCCTGGCAGCACAGG

SEO ID NO: 716

Human mRNA for dihydropteridine reductase (hDHPR)

- 25 TGCTGTCACCAGCTCCTGGTATGATCGGGTACGGCATGGCCAAGGG
 25 TGCTGTTCACCAGCTCTGCCAGAGCCTGGCTGGCAAGAACAGCGGCATGCCGCC
 CGGGGCAGCCGCCATCGCTGTGCTCCCGGTTACCCTGGATACCCCGATGAACAGG
 AAATCAATGCCTGAGGCTGACTTCAGCTCCTGGACACCCTTAGAATTCCTAGTTG
 AAACTTTCCATGACTGGATCACAGGGAAAAACCGACCGAGCTCAGGAAGCCTAA
 TCCAGGTGGTAACCACAGAAGGAAGGACGGAACTCACCCCAGCATATTTTTAGG

INCOME TO A GEORGE TRACEA AGE A TOTE A AGGA AGGA GEORGE CONTRACTOR AGE TO A CONTRACTOR

- 30 CCTCATCTCAGTGCCTATGAGGGGCCTGCCAGAAAAGTCACTAACCTGTCTCAGT GTGGCCTTGTCCAGCCTTGTGTTTTCTGTAACCCCTGTTTGTGGTACGAGATAATG AGTCCTATTTTCTCTCACATAATATGCATTTGCTCTCCTAGGACAGTGTAATACA TTTATGTGAAGTAAAGACATGCGAGACTGGTGGCCTGCAAATAGCATCCGTCAAT CTGTGTTAACTGCATAGGGAGGGCTCTGCATAGCACCTGCTATAGCGGTGTCATG
- 35 TTGGATCGCTTTTGTGACTGTTCATCTGTCCTTGACAGTGGCTGTCATCTTGACTA CTTTGTTGATTTGTTGGTATTGGGGACATTTTAAAGGCTGAGTTATTTTTGAATGT CATGTTTATGTCATAGACGTAGTTTTCGCATCCTTGAATTAAACTGCCTTAACTCC TTTTGTGGTAT
- SEQ ID NO: 717

 aa24g12.r1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:814246 5' similar to gb:D00762 PROTEASOME COMPONENT C8 (HUMAN);, mRNA sequence gi|2191760|gb|AA465593.1|AA465593[2191760]
- CGATGACTCAATCGGCACTGGGTATGACCTGTCAGCCTCTACATTCTCTCTGAC

 45 GGAAGAGTTTTTCAAGTTGAATATGCTATGAAGGCTGTGGAAAAATAGTAGTACA
 GCTATTGGAATCAGATGCAAAGATGGTGTTGTCTTTGGGGTAGAAAAATTAGTCC
 TTTCTAAACTTTATGAAGAAGGTTCCAACAAAAGACTTTTTAATGTTGATCGGCA
 TGTTGGAATGGCAGTAGCAGGTTTGTTGGCAGATGCTCGTTCTTTAGCAGACATA
 GCAAGAGAAGAAGATCCAACTTCAGATCTAACTTTGGCTACAACATTCCACTAA

AACATCTTGCAGACAGAGTGGCCATGTATGTGCATGCATATACACTCTACAGTGC TGTTAGACCTTTTGGGCTGCAGTTTCA

SEQ ID NO: 718

5 zx10e07.s1 Soares_total_fetus_Nb2HF8_9w Homo sapiens cDNA clone IMAGE:786084 3', mRNA sequence gi|2162337|gb|AA448667.1|AA448667[2162337]
ATAAATCTATAGTTTTATTAAGACAAAAACTGACAATGTAGTATGAAGTTTACAT TTAAA

CAAAGTTTACACAGGAATCTAACACATGCCTAAAAGAATTTTACAACGTAGCTCT
AGATGCAAGTCTAGACAATATCAAGAACTGATGGTTCTCATGACTCAAGACAGA
GCATTTTGGGTATGTTACTTATTAGGATTTCTTAAAAAAATTGTTTTTGTGTGTAT
GTGTGTGTTTTAAAAGTGAACCACTGCCCAATATGAAAAGTTTAATCTTCTCCTGAG
ACCAAGGCTTTTGAAATCACTAAACTCTTGGATCAATTCAGTGAAACTTGTGCTG
TCAGTGACTGAACCCTGCCAACAATGGTTTCAGTGTTCAAAGCTCAAAGAAAAC

15 GGCT

SEQ ID NO: 719

Human hyaluronate receptor (CD44) gene, exon 1 gi|180127|gb|M69215.1|HUMSCG01[180127]

- - 25 GAGCATGTGTGGAGAGAGGTGCCCATTCACACTGGCTTGAACACATGGGTTA GCTGAGCCAAATGCCAGCCCTATGACAGGCCATCAGTAGCTTTCCCTGAGCTGTT CTGCCAAGAAGCTAAAATTCATTCAAGCCATGTGGACTTGTTATTGAGGGGAAA AAGAATGAGCTCTCCCTCTTTCCACTTGGAAGATTCACCAACTCCCCACCCCTCA CTCCCCACTGTGGGCACGGAGGCACTGCGCCACCCAGGGCAAGACCTCGCCCTCT
 - 30 CTCCAGCTCCTCTCCCAGGATATCCAACATCCCTGTGAAACCAGAGATCTTGCTC CAGCCGGATTCAGAGAAATTTAGCGGGAAAGGAGAGGCCAAAGGCTGAACCCA ATGGTGCAAGGTTTACGGTTCGGTCATCCTCTGTCCTGACGCCGCGGGGCCAGC GGGAGAAGAAAGCCAGTGCGTCTCTGGGCGCAGGGGCCAGTGGGGCTCGGAGG CACAGGCACCCCGCGACACTCCAGGTTCCCCGA

 - 40 CCAGGGATCCTCCAGCTCCTTTCGCCCGCGCCCTCCGTTCGCTCCGGACACCATG GACAAGTTTTGGTGGCACGCAGCCTGGGGACTCTGCCTCGTGCCGCTGAGCCTGG CGCAGATCGGTGAGTGCCCGCCGCAGGCTGGCAGCAAGATGGGTGCGGGGTGC TCAGCGCGGAC
 - 45 SEQ ID NO: 720

yi63g06.r1 Soares placenta Nb2HP Homo sapiens cDNA clone IMAGE:143962 5', mRNA sequence gi|851402|gb|R76770.1|R76770[851402] AATTCGGAACGAGGNCTGTACAACACAGTGTCATACAGGGATAATGCTATCATA

TTTAATATGAAACAGTGTTACGGGCACAAATTACCCATTTCTACAAAATAAGTGT

GCAAGTGATGCCACATATTATCCATATTCAACTGAGCTGTCATCAAAATACATTT TATTTACAATATGTACTATGATCAGTTGGATATTAAGTTCTAAAATGATTTACTTC ACTGCTACATTATAAAGGTAAAAGCAATGTGTAGGAAAAAGTGTGAGATTGTGT TTTTACATACTGCTTTTGTAGTTGCCATCGCTGGTTCAGTTCGACTTATAACATAT GTCTTGCTTGTAGGATTTAACACCTCCAATAGGGGATTCTTCTAACATTACAGGA GGATTCTTAGGGGATCCGGGGCTTTTTCANCAGTATAT

SEO ID NO: 721

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SEQ ID NO: 722

CC

- Homo sapiens P2U nucleotide receptor mRNA, complete cds gi|984506|gb|U07225.1|HSU07225[984506]
- CGGCACGAGGCACCECGAGAGGAGAAGCGCAGCGCAGTGGCGAGAGGAGCCCC

 TTTGTGGCAGCAGCACTACCTGCCGAGAAAAATGCTGGAGGCTGGGCGTGGCCCC

 AGGCCTGGGGACCTGTTTTTCCTGTTTTCCCGCAGAGTTCCCTGCAGCCCGGTCGA

 CACCTGGAGAGACCAGGGCTCGTCAGGGCGAACCCGTGCAGACCCCTGGA
 - 25 GGTCCAGGCGTGTGCATTCATGAGTGAGGAACCCGTGCAGGCGCTGAGCATCCT GACCTGGAGAGCAGGGCTGGTCAGGGCGATGGCAGCAGACCTGGGCCCCTGGA ATGACACCATCAATGGCACCTGGGATGGGGATGAGCTGGGCTACAGGTGCCGCT TCAACGAGGACTTCAAGTACGTGCTGCCTGTGTCCTACGGCGTGGTGTGCGT GCTTGGGCTGTGTCTGAACGCCGTGGCGCTCTACATCTTCTTGTGCCGCCTCAAG

 - 35 GCCGTGTGGGTGTTGGTGCTGGCCTGCCAGGCCCCCGTGCTCTACTTTGTCACCA CCAGCGCGCGCGGGGCCCGCGTAACCTGCCACGACACCTCGGCACCCGAGCTCT TCAGCCGCTTCGTGGCCTACAGCTCAGTCATGCTGGGCCTGCTCTTCGCGGTGCC CTTTGCCGTCATCCTTGTCTGTTACGTGCTCATGGCTCGGCGACTGCTAAAGCCAG CCTACGGGACCTCGGGCGCCTCCCTAGGGCCAAGCCCAAGTCCGTGCGCACCA
 - 40 TCGCCGTGGTGCTGTCTTCGCCCTCTGCTTCCTGCCATTCCACGTCACCCGC ACCCTCACTACTCCTTCCGCTCGCTGGACCTCAGCTGCCACACCCTCAACGCCAT CAACATGGCCTACAAGGTTACCCGGCCGCTGGCCAGTGCTAACAGTTGCCTTGAC CCCGTGCTCTACTTCCTGGCTGGGCAGAGGCTCGTACGCTTTGCCCGAGATGCCA AGCCACCCACTGGCCCAGCCCTGCCACCCCGGCTCGCCAGGCTGGGCCTGCG
 - 45 CAGATCCGACAGAACTGACATGCAGAGGATAGGAGATGTTTTGGGCAGCAGTGA GGACTTCAGGCGGACAGAGTCCACGCCGGCTGGTAGCGAGAACACTAAGGACAT TCGGCTGTAGGAGCAGAACACTTCAGCCTGTGCAGGTTTATATTGGGAAGCTGTA GAGGACCAGGACTTGTGCAGACGCCACAGTCTCCCCAGATATGGACCATCAGTG ACTCATGCTGGATGACCCCATGCTCCGTCATTTGACAGGGGCTCAGGATATTCAC

10 AA

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SEQ ID NO: 723

aa50e04.s1 NCI_CGAP_GCB1 Homo sapiens cDNA clone IMAGE:824382 3', mRNA sequence

- 20 AACTTGAGCCAAGGGATAAATATAAGCAACCAATGGGCTGCAGGATAGTTGTAC
 AAAGTGTATCATGTATCTTCATAGCTTCTTTTGCCCATATAATGCATTCCACACTTA
 AGTTTCTCCTTCTAAAAGGGGACACGACAAGTTAATATGTCTCATAAATGTCTTA
 AATAAGTTGCATTTCATGGCAAGCCCTCCACTGCCAGCAATGGATATACTCACAC
 CTATTGGAAAAAATCTAAAGTTAACAAACTGGTTTAGTATGGAAATGGTCTATTT
 - 25 GTTCCTCAGCTATGTTTCTGTATCCTACATTAGTGGCTCTCAGGAGG

SEO ID NO: 724

HUMHBC4799 Human pancreatic islet Homo sapiens cDNA similar to alpha-1 antichymotrypsin, mRNA sequence gi|1262485|dbj|D83812.1|D83812[1262485]

- 35 CCCANAGACCCTGAAGCGGTGGAGAGACTCTCTGGAGTTCANAGAGATAGGTGA GCTCTACCTGCCAAAGTTTTCCANCTCGAGGGACTATAACCTGAACGACATNCTT CTCCAGCTGGGCATTGAGGAAGCCTTC

SEQ ID NO: 725

zx84c12.s1 Soares ovary tumor NbHOT Homo sapiens cDNA clone IMAGE:810454 3', mRNA sequence gi|2179839|gb|AA457119.1|AA457119[2179839]
 CTCATCAAAACATGATTTATTAATTTTAAGCAAGAGTAAGCATATGTGATAGTGG CCAGCTTGGGGATGATAGTCTCCTGGTTGATGCACAGTTCAGCACCTGTTGGGT CTTGGCTGTTGGGATGATAATTCTTTTGGGTGAGGGGAACAGCCGTGGTCAAGGC
 TGCCTGCACCCCCATCCAGGCACAGGACCCTGGGCAAAGTCTCAAAAGAGGTAG TGTTTTTACTTTCGCACCAACAATACAACATAAGTATTGGGTACAAAAGAGGAGA TTTCCTTCCCCTCTACCTCAACGGGCAAAAGGCCTTCCATCTTCAGAAGAGGCTT GTGAGGACCATCGGTTGGATGACCTCCTAGTGAGTTCTGGCTCCCATTCAGAAGACA

CAGAGAAACCCACAAAAGGGGCCTGTGGATCTGGTTCCAGGTCTCAAGGGTACA GCTTGGTTACATCCCCAGGCCCC

SEQ ID NO: 726

15

SEQ ID NO: 727

- yr38g10.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:207618 3' similar to gb:L24038_rna1 A-RAF PROTO-ONCOGENE SERINE/THREONINE-PROTEIN KINASE (HUMAN);, mRNA sequence
- 20 gi|1012590|gb|H59758.1|H59758[1012590]
- 25 CTCCCAAAATTTAGAAGTATCCCCAAAGCCAAGAGAAAACCAAATGATGGGAGG AGACAGGGGCTCAGTCTTTGGGCGGGGGTCCCCCAATTTCCAGAAGAACTGGG AAAAGGCACATGGGGGNCCCCCTTCATCTTCCCGGGGTGGGGGAATGGGGGGAT TCCTNAGGGCAGCNTCAGGGGCAGAGACGAACTTTGTTTGGGTTGGTNGGGCAA GGTTCCTTGGGCTTNGGAG

30

SEO ID NO: 728

- Human thyroid hormone receptor alpha 1 (TR-alpha-1) gene, complete cds gi|339662|gb|M24748.1|HUMTHRA1A[339662]
- 40 TCCATCCCACCTATTCCTGCAAATATGACAGCTGCTGTGTCATTGACAAGATCAC CCGCAATCAGTGCCAGCTGTGCCGCTTCAAGAAGTGCATCGCCGTGGGCATGGCC ATGGACTTGGTTCTAGATGACTCGAAGCGGGTGGCCAAGCGTAAGCTGATTGAG CAGAACCGGGAGCGGCGGCGGAAGGAGGAGATGATCCGATCACTGCAGCAGCG ACCAGAGCCCACTCCTGAAGAGTGGGATCTGATCCACATTGCCACAGAGCCCA

CTACGACCCTGAGAGCGACACCCTGACGCTGAGTGGGGAGATGGCTGTCAAGCG GGAGCAGCTCAAGAATGGCGGCCTGGGCGTAGTCTCCGACGCCATCTTTGAACT GGGCAAGTCACTCTCTGCCTTTAACCTGGATGACACGGAAGTGGCTCTGCTGCAG 5 AGAAGAGTCAGGAGCGTACCTGCTGGCGTTCGAGCACTACGTCAACCACCGCA CATGATCGGGGCCTGCCACGCCAGCCGCTTCCTCCACATGAAAGTCGAGTGCCCC ACCGAACTCTTCCCCCACTCTTCCTCGAGGTCTTTGAGGATCAGGAAGTCTAAA GCCTCAGGCGGCCAGAGGGTGTGCGGAGCTGGTGGGGAGGAGCCTGGAGAGAA 10 CGTCCTTGGATAGATTCAGCTCCCACACACACCCCGCACTGCCCAGGTCCCTC CTCAGACCTCCAGCCCTGGGACAGGGCAAACAACTGAACTTGCTATGGAAAGGA CAGTGTGGGAGCTGGGGAGCTGTGTCCTGCAGTTCCCAGGACCCCATCCTCTC AGAAGGTAGGGGAAGGCGGGGAGGATTGAGAAGGGACAAGCCACCTTGACCGT 15 AGGGGAAGGAGGAATGTGGGCTGGGGGAAGATGCCCTCAACTCACCCCCTCACA CACATGAGAGAGAGCCCCCACCCAGTTCCTTGGCCTAGGTCTCCCCTCCAGGCTG AGGGCCTCTCTACTTCCCCAGATGCCTGGGTGCAAAGAACGGCTTGGCTTGGCTC CTCCTCTGGAGGTTAAAATTTATAGTCATTCTAACTGCACTTGGAAACCAAGCAA GGGGAGAAGACAAATGAAGAAAAACT

20

SEQ ID NO: 729

35

SEQ ID NO: 730

CCCCACC

- yr86d03.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:212165 3' similar to gb:Z22548 THIOL-SPECIFIC ANTIOXIDANT PROTEIN (HUMAN);, mRNA sequence gi|1030355|gb|H68845.1|H68845[1030355]
- 40 TTCCTAATACTTTATTGGNTACCTCTAGGCCTGTGTGCGGCTGGGTGGGCTTGG
 GGGAGGGCGTCACTATTCAGCTTCTAGGTGGAGGCATGAGAAGGCCTTGGCTAG
 NCCCTCCAGGGTCCCATACTGTGGAGTTTGGAGGGCAGGTCTGGCCTTTCCTGG
 GTCAGCATAGGGCACCCAGGTNGGGGCACAGGTGGACACCCAGCACAGGCACCT
 AGGCAGGGCACAAGCTCACTATCCGTTAATCGTGTCACCAGGCTTCCAGCCA
- 45 ATTCCTTGCTGTCATCCACGTTGGGCTTAATCGTGTCACTACCAGGCTTCCAGCCA GCGGGANAAACTTTCCCCATGCTCGTCTGTGTACTGGGAAGGNCTGGGACCAGC CGCAGAGCCTANATTCCACGGAGCGTCCCACAGGCAAAT

SEQ ID NO: 731

ab23b05.r1 Stratagene lung (#937210) Homo sapiens cDNA clone IMAGE:841617 5' similar to TR:E183625 E183625 ORNITHINE DECARBOXYLASE ANTIZYME;, mRNA sequence

SEQ ID NO: 732

Human elastase III B mRNA, complete cds, clone pCL1E3

- gi|607029|gb|M18692.1|HUMELA3A[607029]
 CCTATCATCGCAAAACTCATGATGCTCCGGCTGCTCAGTTCCCTCCTTGTGGC
 CGTTGCCTCAGGCTATGGCCCACCTTCCTCTCGCCCTTCCAGCCGCGTTGTCAATG
 GTGAGGATGCGGTCCCCTACAGCTGGCCCTGGCAGGTTTCCCTGCAGTATGAGAA
 AAGCGGAAGCTTCTACCACACCTGTGGCGGTAGCCTCATCGCCCCCGACTGGGTT
 GTGACTGCCGGCCACTGCATCTCGAGCTCCCGGACCTACCAGGTGGTGTTGGGCG

SEQ ID NO: 733

SEO ID NO: 734

45

yv19b06.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:243155 3', mRNA sequence gi|1102102|gb|H94469.1|H94469[1102102] GCAAAACAACATTTATTCTTTTAAAAAAATCTATATACATTGCCATACAAAGATAC

CACATTGAAGCAGTTCTCAGGAACCTTCCAGTGAGCCTTCTCTTATAATTGCCCG AGCAAGATTTCGTGCCAGAGAAAGTCTCAGCATTTCCACCTTGGTGTNCTCTATG TCATCATCCTGGAGCTGCTCGGTATCAGATTCTCCATGCACAGGTCTTCTTGACGT CAAGTCCTCCAGACACCGCATCAACTCATAAGTCTGTTCTGCTGAGAAAATCACC TGTTTCTGTTCCAAAAAGGGGCAAGGCATCTGTCAGCAGAGTTCATCCCAGAAAGA CCGAAGGGGCAATCCGAGACGTCATCAAGGACAGAAGGA

SEO ID NO: 735

5

aa91g07.s1 Stratagene fetal retina 937202 Homo sapiens cDNA clone IMAGE:838716 3'
 similar to TR:G173234 G173234 RIBOSOMAL 5S RNA-BINDING PROTEIN ;, mRNA sequence

gi|2180364|gb|AA457644.1|AA457644[2180364]

- 15 CCAAACCAAATGGATATCTGCTTTTAAGATTAGAATTTGTTCTTCATCCTTAAAGC AGAACTCATTGAGATGAAAAGATGCTCTTAATTTATCACAGAACTGTGTATTTAA TAGTATGCTTATTAAAATCACGAAGTGTACTGGAATGCTAAGATAAAAGAACTGT ATAGTTTCTGTTATGTAATACGAGAATAGAAATGTTATTAAAATCTTTCTATAATT TCCAGTGCTTCTGTTTTGAAGAACAAAAGGCTTAATCCCCAAGAGGAAGTAGATAT
- 20 GCCAGTGTTTTCTACATTGATCCTGAATTTGCTGAAGATCCA

THE LOSEQUE NO. 736 HOW RESERVED AND THE PROPERTY OF THE PROPERTY AND A SECOND WAS AN ART OF THE VERY

- FOC the Hasapiens CD18 exon 14 gi|29753|emb|X63924.14|HSCD18X14[29753] the Section 14 gi|29753|emb|X63924.14|HSCD18X14[29753]
- CTCCCCGCAGCTCCTGCGCCGAGTGCCTGAAGTTCGAAAAGGGCCCCTTTGGGAA
 - 25 GAACTGCAGCGCGGCGTGTCCGGGCCTGCAGCTGTCGAACAACCCCGTGAAGGG CAGGACCTGCAAGGAGGGACTCAGAGGGCTGCTGGGTGGCCTACACGCTGGA GCAGCAGGACGGATGGACCGCTACCTCATCTATGTGGATGAGAGCCGAGGTGA GGCCGC

30 SEQ ID NO: 737

- ye81h02.s1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:124179 3', mRNA sequence gi|751008|gb|R01272.1|R01272[751008]

- 40 ATATTCCCCTCAGGTTCCCGGTTTCCATTTTGTT

SEQ ID NO: 738

- zx35f11.s1 Soares_total_fetus_Nb2HF8_9w Homo sapiens cDNA clone IMAGE:788493 3', mRNA sequence gi|2166225|gb|AA452556.1|AA452556[2166225]

TGAACACTGAAAAGAACAATATATATACTGTAAAATATGATGAATAAACCAAATG TAGCTATAAGAATCTTAAAGGATGATTATAGAAAAGGGA

SEQ ID NO: 739

15

- **SEO ID NO: 740**
- ye40b03.r1 Soares fetal liver spleen 1NFLS Homo sapiens cDNA clone IMAGE:120173 5', mRNA sequence gi|734317|gb|T95693.1|T95693[734317]
- AGATGTTCCAGGAGTAGAATTCCTGACTGCTGTGTGAAAGTGAACTGCTACTCCA

 20 TCTCTGAAACATATCTGAGAAACGGGGCAGAAAACCAGTGTAAACTGCTCGTGG
 TGAAATTATTGAACATTGAAGTGTGAGGCTTGTCCTAAGAGCACGTCACCTCCCT
 AGACACAGATTCTGCATGTCCTTCCCTGGTAGGGATCCTCCAGTTCCGTTTCTC
 AGGCGAAGTAACCAGAGGTTCCAGTCTGGTNTTGCTTTCTGGGGGAGGAAGGAC
 AGGAGGCACCTAGGTTAATAGGATTCCCAGGTACTTGATTGGGGCACCCACAC
 - 25 ATGGANTTCAGGAGGGGGACCTTAAGGCCNTTCAGGCAGG

SEQ ID NO: 741

Human (clone HSY3RR) neuropeptide Y receptor (NPYR) mRNA, complete cds gi|189313|gb|L01639.1|HUMNYRECA[189313]

- 30 CGCATCTGGAGAACCAGCGGTTACCATGGAGGGGATCAGTATATACACTTCAGA TAACTACACCGAGGAAATGGGCTCAGGGGACTATGACTCCATGAAGGAACCCTG TTTCCGTGAAGAAAATGCTAATTTCAATAAAATCTTCCTGCCCACCATCTACTCC ATCATCTTCTTAACTGGCATTGTGGGCAATGGATTGGTCATCCTGGTCATGGGTT ACCAGAAGAAACTGAGAAGCATGACGGACAAGTACAGGCTGCACCTGTCAGTGG 35 CCGACCTCCTCTTTGTCATCACGCTTCCCTTCTGGGCAGTTGATGCCGTGGCAAAC
- TGGTACTTTGGGAACTTCCTATGCAAGGCAGTCCATGTCATCTACACAGTCAACC TCTACAGCAGTGTCCTCATCCTGGCCTTCATCAGTCTGGACCGCTACCTGGCCATC GTCCACGCCACCAACAGTCAGAGGCCAAGGAAGCTGTTGGCTGAAAAGGTGGTC TATGTTGGCGTCTGGATCCCTGCCCTCCTGCTGACTATTCCCGACTTCATCTTTGC
- 45 AAATCATCAAGCAAGGGTGTGAGTTTGAGAACACTGTGCACAAGTGGATTTCCA
 TCACCGAGGCCCTAGCTTTCTCCACTGTTGTCTGAACCCCATCCTCTATGCTTTC
 CTTGGAGCCAAATTTAAAACCTCTGCCCAGCACGCACTCACCTCTGTGAGCAGAG
 GGTCCAGCCTCAAGATCCTCTCCAAAGGAAAGCGAGGTGGACATTCATCTGTTTC
 CACTGAGTCTGAGTCTTCAAGTTTTCACTCCAGCTAACACAGATGTAAAAGACTT

TTTTTTTATACGATAAATAACTTTTTTTTAAGTTACACATTTTTCAGATATAAAAG ACTGACCAATATTGTACAGTTTTTATTGCTTGTTTGGATTTTTGTCTTGTGTTTCTTT AGTTTTTGTG

GTGGGCAGCCTGCTTCCTGCATGGAACTGGGAGCTGTATGGAGTTGACGACAA

5 SEQ ID NO: 742 >AA504554 CACCACGGTGACCGTTTTCATCAGCAGCTCCCTCAACACCTTCCGCTCCGAGAA GCGATACAGCCGCAGCCTCACCATCGCTGAGTTCAAGTGTAAACTGGAGTTGCTG

10 GTTCTACAGCAAGCTG GATCAAGAGGATGCGCTCCTGGGCTCCTACCCTGTAGATGACGGCTG

SEQ ID NO: 743 >M11723

- 25 ACCGCCTGTGCCACTGCCCGGTGGGCTACACCGGACCCTTCTGCGACGTGGACAC
 CAAGGCAAGCTGCTATGATGGCCGCGGGCTCAGCTACCGCGGCCTGGCCAGGAC
 CACGCTCTCGGGTGCGCCCTGTCAGCCGTGGGCCTCGGAGGCCACCTACCGGAAC
 GTGACTGCCGAGCAAGCGCGGAACTGGGGACTGGGCGCCACGCCTTCTGCCGG
 AACCCGGACAACGACATCCGCCCGTGGTGCTTCTGCTGAACCGCGACCGGCTG
- 30 AGCTGGGAGTACTGCGACCTGGCACAGTGCCAGACCCCAACCCAGGCGCGCCT CCGACCCCGGTGTCCCCTAGGCTTCATGTCCCACTCATGCCCGCGCAGCCGGCAC CGCCGAAGCCTCAGCCCACGACCCGGACCCCGTCTCAGTCCCAGACCCCGGGAG CCTTGCCGGCGAAGCGGGAGCAGCCGCCTTCCCTGACCAGGAACGGCCCACTGA GCTGCGGGCAGCGGCTCCGCAAGAGTCTGTCTTCGATGACCCGCGTCGTTGGCGG
- 35 GCTGGTGGCGCTACGCGGGGCCACCCCTACATCGCCGCGCTGTACTGGGGCCA CAGTTTCTGCGCCGGCAGCCTCATCGCCCCCTGCTGGGTGCTGACGGCCGCTCAC TGCCTGCAGGACCGGCCCGCACCCGAGGATCTGACGGTGGTGCTCGGCCAGGAA CGCCGTAACCACAGCTGTGAGCCGTGCCAGACGTTGGCCGTGCGCTCCTACCGCT TGCACGAGGCCTTCTCGCCCGTCAGCTACCAGCACGACCTGGCTCTGTTGCGCCT
- 40 TCAGGAGGATGCGGACGGCAGCTGCGCGCTCCTGTCGCCTTACGTTCAGCCGGTG
 TGCCTGCCAAGCGGCGCGCGCGCGCGCCCTCCGAGACCACGCTCTGCCAGGTGGCC
 GGCTGGGGCCACCAGTTCGAGGGGGGGGGAGGAATATGCCAGCTTCCTGCAGGAG
 GCGCAGGTACCGTTCCTCTCCCTGGAGCGCTGCTCAGCCCCGGACGTGCACGAT
 CCTCCATCCTCCCCGGCATGCTCTCTCTCAAGAAACGTAAACGCAAACGCA
- 45 GTGCCAGGGTGATTCCGGAGGCCCGCTGGTGTGTGAGGACCAAGCTGCAGAGCG CCGGCTCACCCTGCAAGGCATCATCAGCTGGGATCGGGCTGTGACCGCAA CAAGCCAGGCGTCTACACCGATGTGGCCTACTACCTGGCCTGGATCCGGAGCA CACCGTTTCCTGATTGCTCAGGGACTCATCTTTCCCTCCTTGGTGATTCCGCAGTG

AGAGAGTGGCTGGGGCATGGAAGGCAAGATTGTGTCCCATTCCCCAGTGCGGCCAGCTCCGCGCCAGGATGGCGCAGGAACTCAATAAAGTGCTTTGAAAATGCTG

SEQ ID NO: 744

5 >S60489

CTACTCCTAGATATTTGGCATGATCTTCAGTATGATCTTGTGCTGTGCTATCCGCA GGAACCGCGAGATGGTCTAGA

SEQ ID NO: 745

10 >M59916

15 TGGCGCTGGCGCTGGCTCTGTCTGACTCTCGGGTTCTCTGGGCTCCGGC
AGAGGCTCACCCTCTTTCTCCCCAAGGCCATCCTGCCAGGTTACATCGCATAGTG
CCCCGGCTCCGAGATGTCTTTGGGTGGGGGAACCTCACCTGCCCAATCTGCAAAG
GTCTATTCACCGCCATCAACCTCGGGCTGAAGAAGGAACCCAATGTGGCTCGCGT
GGGCTCCGTGGCCATCAAGCTGTGCAATCTGCTGAAGATAGCACCACCTGCCGTG
20 TGCCAATCCATTGTCCACCTCTTTGAGGATGACATGGTGGAGGTGTGGAGACGCT

20 TGCCAATCCATTGTCCACCTCTTTGAGGATGACATGGTGGAGGTGTGGAGACGCT
CAGTGCTGAGCCCATCTGAGGCCTGTGGCCTGCTCCTGGGCTCCACCTGTGGGCA
CTGCGGACATTTCTCATCTTGGAACATCTCTTTGCCTACTGCCGAAGCCGCCCGAGCCCCAGGTGCCCCTGTCAGCCGCATCCTCTCCT
CACTGACCTGCACTGGGATCATGACTACCTGGAGGGCACCGGACCCTGACTGTGC

45 GCCGTCTGGCTACTCTTTGTGCCCAGCTCTCTGCCCGTGCTGACAGCCCTGCTCTG
TGCCGCCACCTGATGCCAGATGGGAGCCTCCCAGAGGCCCAGAGCCTGTGGCCA
AGGCCACTGTTTTGCTAGGGCCCCAGGGCCCACATTTGGGAAAGTTCTTGATGTA
GGAAAGGGTGAAAAAGCCCAAATGCTGCTGTGGTTCAACCAGGCAAGATCATCC
GGTGAAAGAACCAGTCCCTGGGCCCCAAGGATGCCGGGGAAACAGGACCTTCTC

SEQ ID NO: 746

>W74362

5

TGAAGATGGAGCTAATCTTTCCTCTGCTCGTGGCATTTTGTCGCTTATCCAGTCTT

10 CTACTCGTAGGGCATACCAGCAGATCTTGGATGTGCTGGATGAAAATCGCAGAC
CTGTGTTGCGTGGTGGGTCTGCCGCCACTTCTAATCCTCATCATGACAACGT
NAGGTATGGCATTTCAAATATAGATACAACCATTGAAGGAAAGACCCCCNCNCC
NCGACTGTNNTAGATGCANCN
CCCCCCCAGAAGACAGATAATCAAACTAAATAGACGTCTA

15

SEQ ID NO: 747 >N71365

SEQ ID NO: 748 >AA454662

SEQ ID NO: 749 >AA450180

SEQ ID NO: 750 >N76338

GCGANTGGCATTGAGCTACAGGCAGGAGATGAGGTGGAGTTCTCAGTGATTCTT
AATCAGCGCACTGGCAAGTGAGCGCCCTGTAATGTTTGGCGAGTCTGTGAGGGC

5 CCCAAGGCTGTTGCAGCTCCTCGACCTGATCGGTTGGTCAATCGCTTGAAGAATA
TCACTCTGGATGATGCCAGTGCTCCTCGCNTAANANGNGNTTCTTCGTCAGCCAN
GGGGACCAGATAACTCAATGGGTTTTGGTGCAGAAAGAACATCCGTCAAGCTG
GTGTCATTGACTAACCACNTCCACAANGCACACCATTTAATCCACTATGATCAAG
TTGGGGGAATACTTATTCTATTGGNGCTATTACACCAGTTTTAAANACCTTCCNCG

10 TCTGGGAATACTTATTCTATTGGNGCTATTACACCAGTTTTAAANACCTTCCNCG GGGTTATGGTTTTAAAAAAAATAAATTTTAGAAAACCCTT TTAAATAATGCACAGTTGCAGCCTGGNAAAA

SEQ ID NO: 751

>M60626

15

CCCAGAGCAAGACCACAGCTGGTGAACAGTCCAGGAGCAGACAAGATGGAGAC AAATTCCTCTCCCCACGAACATCTCTGGAGGGACACCTGCTGTATCTGCTGGC TATCTCTTCCTGGATATCATCACTTATCTGGTATTTGCAGTCACCTTTGTCCTCGG 20 AGTCACCACCATCAGTTACCTGAACCTGGCCGTGGCTGACTTCTGTTTCACCTCC ACTTTGCCATTCTTCATGGTCAGGAAGGCCATGGGAGGACATTGGCCTTTCGGCT YEAR OF CONTROL OF THE STREET 25 @GGTGATGGCTCTGCTCCTCACATTGCCAGTTATCATTCGTGTGACTACAGTACCTG GTAAAACGGGGACAGTAGCCTGCACTTTTAACTTTTCGCCCTGGACCAACGACCC TAAAGAGAGGATAAATGTGGCCGTTGCCATGTTGACGGTGAGAGGCATCATCCG GTTCATCATTGGCTTCAGCGCACCCATGTCCATCGTTGCTGTCAGTTATGGGCTTA TTGCCACCAAGATCCACAAGCAAGCTTGATTAAGTCCAGTCGTCCCTTACGGGT CCTCTCCTTTGTCGCAGCAGCCTTTTTTCTCTGCTGGTCCCCATATCAGGTGGTGG 30 CCCTTATAGCCACAGTCAGAATCCGTGAGTTATTGCAAGGCATGTACAAAGAAAT TGGTATTGCAGTGGATGTGACAAGTGCCCTGGCCTTCTTCAACAGCTGCCTCAAC CCCATGCTCTATGTCTTCATGGGCCAGGACTTCCGGGAGAGGCTGATCCACGCCC TTCCCGCCAGTCTGGAGAGGGCCCTGACCGAGGACTCAACCCAAACCAGTGACA 35 CAGCTACCAATTCTACTTTACCTTCTGCAGAGGTGGCGTTACAGGCAAAGTGAGG AGGGAGCTGGGGGACACTTTCGAGCTCCCAGCTCCAGCTTCGTCTCACCTTGAGT TAGGCTGAGCACAGGCATTTCCTGCTTATTTTAGGATTACCCACTCATCAGAAAA AAAAAAAAGCCTTTGTGTCCCCTGATTTGGGGAGAATAAACAGATATGAGTTT ATTATTGACTTCTTTTTTGATTTTGGACCTCAGCCTCGGGTGGTCAGGGTGGGAAA 40 TGATAGGAAGAAGCTGTCATCTGCATCCTAGTTTGCCTGAAATGAACCCAAATAA TACCCATTATTATTAGTCCTGAATTATGAGTAGTGAATGATACCCATCATTCTGGC ATCATGATGAGTAGTGTCCACTTCCATTCTGAAAAGTGCCCTGCTGTGAAAAATA AATTATATAGTCATCCTAGGTAAATGAAGGAGGAGGGGAGAAGTGTGAAAGAGTA TGGCTTAAATCAGACAAGATATACAAGAAGATACTTTATATAGGGCAGGAGCGG 45 GAGGTCAGGAATTCGAGAACAGCCTGGCCAACATGGTGAAACCCTGTCTCTACT AAAAATACAAAAATTAGCTGGGCGTAGTGGCAGGCTCCCGTAATCCCAGCTACT CAGGAGACCGAGGCAGGAGATCGCTTGGACCTGGAAGGCGGAGGTTGTAGTGA

SEQ ID NO: 752

5 >X70070 TCAAGCTCGCCCGCGCAGCCGAGCCGGGCTGGGCGCTGTCCTCGGGGGCCTG GGGAACCGCGGTTTGGAGATCGGAGGCACCTGGAACCCGTGGCAAGCGCCGA GCCGGGAGACAGCCGAGGAACCACGGGTTCTGGAGCTAGGAGCCGGAAGCTG GGAGTCCGGAGGAGCCGGAGCCCGGAGCCCGGGGCGCGCGTCTG 10 GGTCTGGCGCTTCCCGACTGGACGGCGCCCCGCTGGTCTTCGCCACGCGCCCTC CCCTGGGCTCGCGTTCATCGGTCCCCGCCTGAGACGCGCCCACTCCTGCCCGGAC TTCCAGCCCGGAGGCGCCGGACAGAGCCGCGGACTCCAGCGCCCACCATGCGC CTCAACAGCTCCGCGCGGGAACCCCGGGCACGCCGGCCGACCCCTTCCAG CGGGCGCAGGCCGGACTGGAGGAGGCGCTGCTGGCCCCGGGCTTCGGCAACGCT 15 TCGGGCAACGCGTCGGAGCGCGTCCTGGCGGCACCCAGCAGCGAGCTGGACGTG AACACCGACATCTACTCCAAAGTGCTGGTGACCGCCGTGTACCTGGCGCTCTTCG TGGTGGCACGGTGGCAACACGGTGACGGCGTTCACGCTGGCGCGGAAGAAGT CGCTGCAGAGCCTGCAGAGCACGGTGCATTACCACCTGGGCAGCCTGGCGCTGT CCGACCTGCTCACCCTGCTGGCCATGCCCGTGGAGCTGTACAACTTCATCTG 20 GGTGCACCACCCCTGGGCCTTCGGCGACGCCGGCTGCCGCGGCTACTACTTCCTG CGCGACGCCTGCACCTACGCCACGGCCTCAACGTGGCCAGCCTGAGTGTGGAG NO CONTROL OF THE PROPERTY OF

GCTCAGGCCTCAAGATCTTCAGCTGTGGCCTCTCGGGCTCGGCAGAAGG GACGCCGGATCAGGGCCTGGTCTCCAGCACCTGCCCGAGTGGCCGTGGCCAGG ATGGGGTGCGCATTCCGTGTGCTTTGCTTGTAGCTGTGCAGGCTGAGGTCTGGAG CCAGGCCCAGAGCTGGCTTCAGGGTGGGGCCTTGAGAAGGGGAATGTGGGACAG 5 GGGCGATGGTGCCTGGTCTCTGAGTAAGATGCCAGGTCCCAGGAACTCAGGCTTC AGGTGAGAAGGAGCGGTGTGTCCAGGCACCGCTGGCCGGCAGCCCTGGGCTGAG GCACAGACTCATTTGTCACCTTCTGGCGGCGGCAGCCCTGGCCCCGGCCTCCAAG CAGTTGAAAAAGCTGGCGCCTCCTTGGTCTCTAGGATCCAGGCTCCACAGAGCAC ATGACTAGCCAGGCCCCTGGCTTAAGAAGGTCGCCTAAGCCTAAGAGAAGACAG 10 TCCCAGGAGAAGCTGGCCGGGACCAGCCAGGAGCTGGGAGCCACAGGAAGCAA AAGTCAGCCTTTCCTCAAGGGATTTCCCTGTCTCAGAGCAGCCTTTGCCCCAGG GAAATGGGCTCTGGGCTGCCTGCACCGGCCATGTCGACCCAGGACCCGGA CACCTGGTCTTGGGCTGTTCAGCCACTTTGCCTTCTCTGGACTCAGTTTCCCCG TCTGAGAAATGAGAGTCGAATGCTACAGTATCTGCAGTCGCTTGGATCTGGCTGT 15 TGAGTTGACGGGTTCCTTGAACCCCACAAAATCCCTCTCCAACCACAGGACCCTT CGGCTCACCAAGAACGGGGCCCAGGGGAGTCAGGCCTATTCGCTGCACTTCCTG CCAAACTTTGCCCCCACAAGCCTGGTCATCAGCCAGGCAGCCCTCCCAGTGCCCA AGGGCCACCAACCCCAGGGAAACAGGGCCAGCACAGAGGGGCCTTCCTCCCCA 20 GATGTCCAGAGGTCGGTGCAGCCCCTATCCCTGCTCAGGAGTGGGCTCAGAGTCT AGCAAATGCTAAGGCCCCTCAGGCTGGGCTCTGAACGAGGACCTGGACTCAGAG I CONTROL OF THE CONT TCAGGATGGTGCTCTGAGAGAGGGCAGAGTGGATGCCCCACTGCCCTAGACCCT CGGTAGACGTGGGGTCTCTGGGGCGGGTCTGTGGCTGTGACTGAAGTCGGCTTT TCCATGCACCACAGACACCCACGACACCTGATCTCGTATCACTAGCTTGCGGC CAGGTCATGATGTGGCCCCGGAAGCTGGCCCTGCGTGCCATGAGTGCGTCGGTCA 30 TGGAGTCCGGAGCCCTGAGCCGCCCTGGTGACGGCACAGCCCTCACAGCTC CTCTCAATAAAGGTGGCCGAAGGGCCTCGATGTGG

SEQ ID NO: 753

35 >X58454 ATGCTGCCGCCAGGCAGCACGGCACCGCGTACCCGGGGCAGTTCGCTCTATAC CAGCAGCTGGCGCAGGGAACGCCGTGGGGGGGCTCGGCGGGGGCACCGCCACTG GGGCCCTCACAGGTGGTCACCGCCTGCCTGCTGACCCTACTCATCATCTGGACCC TGCTGGGCAACGTGCTGGTGCGCAGCCATCGTGCGGAGCCGCCACCTGCGCG 40 CCAACATGACCAACGTCTTCATCGTGTCTCTGGCCGTGTCAGACCTTTTCGTGGC GCTGCTGGTCATGCCCTGGAAGGCAGTCGCCGAGGTGGCCGGTTACTGGCCCTTT GGAGCGTTCTGCGACGTCTGGGTGGCCTTCGACATCATGTGCTCCACTGCCTCCA TCCTGAACCTGTGCGTCATCAGCGTGGACCGCTACTGGGCCATCTCCAGGCCCTT CCGCTACAAGCGCAAGATGACTCAGCGCATGGCCTTGGTCATGGTCGGCCTGGC 45 ATGGACCTTGTCCATCTCATCTCCTTCATTCCGGTCCAGCTCAACTGGCACAGG GACCAGGCGGCCTCTTGGGGCGGGCTGGACCTGCCAAACAACCTGGCCAACTGG ACGCCCTGGGAGGAGCTTTTGGGAGCCCGACGTGAATGCAGAGAACTGTGAC TCCAGCCTGAATCGAACCTACGCCATCTCTTCCTCGCTCATCAGCTTCTACATCCC CGTTGCCATCATGATCGTGACCTACACGCGCATCTACCGCATCGCCCAGGTGCAG

15 SEQ ID NO: 754 >D13538

- - 25 GCCTGGACCGCTACTGGTCGGTGACGCAGGCCGTCGAGTACAACCTGAAGCGCA CACCACGCCGCGTCAAGGCCACCATCGTGGCCGTGTGGCTCATCTCGGCCGTCAT CTCCTTCCCGCCGCTGGTCTCGCTCTACCGCCAGCCCGACGCCGCCGCCTACCCG CAGTGCGGCCTCAACGACGAGACCTGGTACATCCTGTCCTCCTGCATCGGCTCCT TCTTCGCGCCCTCCTCATCATGGGCCTGGTCTACGCGCGCATCTACCGAGTGGC

 - 40 CTTCTGGATCGCTACTGCAACAGCTCGCTCAACCCGGTCATCTACACGGTCTTC AACCAGGATTTCCGGCGATCCTTTAAGCACATCCTCTTCCGACGGAGGAGAAGG GGCTTCAGGCAGTGACTC

SEO ID NO: 755

45 >N76944

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SEQ ID NO: 756

5 >AA451716

10 GCAGCTGTGATCCAGCAGCAGCTGGCAAAGCTTAGTAAGCAACCTCATCCCCAG ATGCATCCGCTCAGCCAGTGTTGTGATTGCTAGATACTATCTGTAAGTGAACCAA ACTAAAATTCATTTATGAACCAAGAAAGGAAGCCAAGTTGAAAAGGTCTCGAGT TAAATCGAGAATGATTCAGGCGGGCCGGCTCTCTGAGCA CCTTTGGATGCACTTCAGCTTCTGTCTTG

15

SEQ ID NO: 757

>H19264

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20 GGTGGTGAACCTGGTNGTGGAGAGCACCCTACTTTAGCCAACTTGGGCAGGGT GGCCCAGGTCCTGAGGCTGATGCGGATCTTCCGATCTTAAAGCTGGCCAGGCACT

AND ACTGGCCTCCGCTCCCT AND MARKET AND AND ACTION OF A STATE OF A ST

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25 SEQ ID NO: 758

>AA598527

30 CCACCATTTGGCTAATATTATTTCATTAAAGACTGAATTTAGATTTTAGGAAATA AAATATGGAATCTGTTATAATGTCCCAATTTATACTACAGTATTAATCTCAATCCT GATCATTACATAATTATAGCATTTACCAATCTGTGATTTTATAAATTAACCAAATT TGTTAAATTAAGAAGAAATTCATAGACACCATTTTTTTTCCTGTTACAACATATGG AAAAGCCATCAAAAAAAACTTAACAGAACCAAATCAAAAAAGAAGTATATTTATGC

35 TAAAGTTACTTTCTGTCCAGGTCGAAACATTGTTC

SEQ ID NO: 759

>AA286908

SEQ ID NO: 760 >AA280924

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10 TGGTCAACACGCGGGCATCGCTTCAAGGTTGCTGATCCCACACCCTTTCATATT CAAGCTGAAGTGACGATGAAAACAAATTTCTTTGGTACCCGAGATGTGTGCACA GAATTACTCCCTCTAATAAAACCCCAAGGGAGAGTGGTGAACGTATCTAG

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15 >AA279601

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20 GGGAGTGCAGTCATCACGGTTGT G

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GTTAAAACATGAAAAAAATTTTATTGTTTTAGACAAAGAGGCCACTTTTGGAAA 25 ATAATACTTTTTTTTTTTTGAATCAGGTGAAGACAGAGTTAAAATCACATA GGATTTGCATTTTAAAAAAGGAAAGCACTAGGATTGTTGGCACTGGAGTAACTA TTTACACTGAACAGAGGTTTGGCCTTTTACATAACATCGATACAATGCATTTTCC AAAGTCTGAGAAATAACAAGGTTCTGTCTCGAATGCTTCACAGAGGAGGTTCGG ATTTGGGGACAAGTGTCATTAATGAGGGCCATGGAAGTTCGTCAGCTTCAGAGTC 30 ACATGCAATCTGATCCTGGGCGGTTCCCCNGCTGGGGAGCACTTGGCTACGGAAT

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SEQ ID NO: 763 >T61575

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45 **SEQ ID NO: 764**

>R23586

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SEQ ID NO: 766

SEQ ID NO: 767 >U39613

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SEQ ID NO: 768

5 >H91337

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SEO ID NO: 769

>M29870

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SEQ ID NO: 772

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SEQ ID NO: 773 >L15189

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SEQ ID NO: 7.7.5

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45 SEQ ID NO: 777

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SEQ ID NO: 779

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SEQ ID NO: 782

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SEQ ID NO: 783

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SEQ ID NO: 785 >AA477082

SEQ ID NO: 786

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SEQ ID NO: 788 >AA401448

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SEQ ID NO: 789

>T84762

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- 15 ATAAATACATGTTTTGGAAATACAGTGACCTCTTGCAGTGTCACAAAAGTGCAAA GTGATATTAGCTGTCATCTGCAATACAGAATCTCATTGCTTTTTGCACATGGAGCA TATAGGGAAACTCCANACAGATCACAATGAGGGTTTCTAAATCTGTTGGGGTTCT GTCTTCTATTGGGGTTCTGTGAAGGCAAACCACTGTAGGCTTTAGCTGGGGTTCN GTCCTATGGACTCGTTGGGGGGNATGCCNTG
- 20 GGTTTTTCCATNCTTACCTGGCAGTCTTGGGGGGGT

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30 AGTNGGTGCCAGGGTGCAAGTTAGGCTAAAGAAGCCACCACTTATTCCTCTCT TGCCCATTTNTGGGGGGGCAAAGGCCATTTGGTCACCCAAGAGTCTTTCCAGGGG GACCCACAGATATTGCCATGTCCCTNCACACGTCTTTGGNGTCCTTAACN

SEQ ID NO: 791

35 >AA424743

45 SEQ ID NO: 792

>AA489331

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CAGCAACATGTTTCTCAGCCGAATTACATTCATAGAAAGTGTCCAGATTCTAAAA TCAAATTAAAAGCATGTAATCCAAAGCCTGAAAAAGCAAACAGCTTTAGGGGCT GACTCCATTAGCGTTCCATAGACTGTGCTTTTAACCGTTCAGTTCATGTTTAATGG CCCATCGGTTCCTTACATACCATCAGCTTATGCTGTGGCCAAAAGAAGTGTTCTT GTGGCTTGGTACTCGTCCCTTCAAACAGTAAACAAGAAAGTGCAGACAGTGCTG CCAGAGACAG

SEQ ID NO: 793

>T67104

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SEQ ID NO: 794

20 >R65792

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- 25 GGCTAAAGCAATTCATCCAAGGGCCACCGGAAGTAATTAGAGCTTTGAAAAAAT CTGTTTGTTCAGGCAGAGAGCTATATTTGGGGGGAAGCATTACAGAACGAAAGA GATCTTTAGGGNACAGTTTTGGGGTNGGCCNGCAAATTTTAGAGGCTATTTNCT AAGGAAGGGNATTTTATTAATATTTGGGTTTTTCCCG
 - 30 SEQ ID NO: 795

>T90621

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- TATGCCTGACACGCCGGAGGGCTNGAGGGGGAACACACTGAAAGCAGTACCAGG GAGCAGTGCATCTCACAGANCCATTTNTTCATGCCATGAAGTAAACGGTACTTAT ACAAGTGTACAGTGACGTTCCACGNTCCCCATCTAACACGGNTTGCTGGAANTTT ACAGGCAGACTGACGTTTTCTTTCACATGTACTCCAAGTAAATCTGGTTAGTGAT GACCNGGGGGCAGGCGCTGAAGCTTTTCAAAAGCCTTACTTCTTTTATCAGCAGCC
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SEQ ID NO: 796

>AA464067

SEQ ID NO: 797 >AA291163

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SEQ ID NO: 798

>N53024

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20 ACTGCATATTTAAAGCATGTGTTCACACTGTGTGTAAACATTCACTGGAAGATTT TTTCNTTGTGCATTGCTGACTGTTCCAACATNACAAGTATTATTAAAATTAAATAT TAACTGGACCGGAAAAAAAGAAAAAAAAAAAGNTCCTTACCCG

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25 >AA398230

> GCGGAATGGGAAGCAGTTTATGGAGTTAAGTGGGGCTCTGCTATTTCCCCCAAGA AGGACTCGGAAGATGTTGATTCCAGGGCAGAGTGAGGGGCAGACGGGATGAGG ${\sf CTCTTCTGTAAAGTCCAACAGACGCTCACAGATGCTGGGAGGCTGGGGACTGCC}$ AGGTTGGGAGCCTCACCAGAGAGCCTCACTGCATTGACCCCACACCACCACTC

1 1 17 17 1

了**我只要只要你的**我们的,这些好,这一个简单是一种的概念的是**说**话,这么

30 ACCCAGCACACAGGGGCCTCTCCTCACGCTCCCAGGCCACCAGGATGGCCCCC AGGTTCACACACAGGCACACGCACACGCTGCACTCACCACGCACTGAAGGGC ATCACAGCCCCAAGTCTGGGTAAGAAATTCTCCAC

SEQ ID NO: 800

35 >H21107

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40 TCATGGCCTTCTTAAACAGCTTTCTTAATCCTTTCTGGGAAATATCCTTTGGTTCA TTTTTATTGCCCCTCTCTNGGGCAAAACAAAGTATGTTAACGCAGGNATCAGTGA GTTATNTCCTAGGCACTTGTAAGGCAATATCCTTACCAAGAGGGACCATTCAACT TTTGTAATAATCCGTNAAGCG

45 **SEQ ID NO: 801**

> zd20g08.s1 Soares fetal heart NbHH19W Homo sapiens cDNA clone IMAGE:341246 3' similar to WP:ZK970.2 CE02402 CLPP-LIKE PROTEASE;, mRNA sequence gi|1365390|gb|W58658.1|W58658[1365390] GCGACCGCCGAGCGACAGATCCAGAACGGCCTGGCCTGCAGCGTGCCTGACGC

GAACGGCANCCCGGGCTCTCCCGCTCATTCCCATCGTGGTGGAGCAGACGGGTC
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TGCGTCATGGGCCCGATCGATGACAGCGTTGCCAGCCTTGTTATCGCACAGCTCC
TCTTCCTGCAATCCNGAGAGCAACAAGAAGCCCATCCACATGTACATCAACAGC
5 CCTGGTGGTGTGGTGACCGCGGGCCTGGCATCTACGACACGATGCAGTACATCCT
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TTCTCGCCGCCGGAACCCCAGGCATGCGCCACTCGCTCCCCAACTCCGTATCAT
GATCCACCAGCCCTCAGGAGGCGCCCG

- 20 TATTTCTCTATTTCATAATCAGTAATAGTGTCATATAAACTCATTTATCTCCTCTT CATGGCATCTTCAATATGAATCTATAAGTAGTAAATCAGAAAGTAACAATCTATG GCTTATTTCTATGACAAATTCAAGAGCTAGAAAAATA

SEQ ID NO: 803

- ab35g03.s1 Stratagene HeLa cell s3 937216 Homo sapiens cDNA clone IMAGE:842836 3' similar to gb:M93056 LEUKOCYTE ELASTASE INHIBITOR (HUMAN);, mRNA sequence gi|2216491|gb|AA486275.1|AA486275[2216491]
 - TCTGGACAGTGGTTTTATTGGTAAAGATATAAGACATATTGGCTCTATTAAAAAC

(19) World Intellectual Property Organization International Bureau





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(71) Applicant: ORTHO-CLINICAL DIAGNOSTICS, INC. [US/US]; 1001 US Highway 202, Raritan, NJ 08869 (US).

- (72) Inventors: WAN, Jackson; 10929 Corte Luz Del Sol, San Diego, CA 92130 (US). WANG, Yixin; 12511 El Camino Real, Unit E, San Diego, CA 92130 (US).
- (74) Agents: PELTO, Don et al.; McKenna & Cuneo LLP, 1900 K Street NW, Washington, DC 20006 (US).

(81) Designated States (national): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZM, ZW.

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For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

(54) Title: EXPRESSION PROFILES AND METHODS OF USE

(57) Abstract: The present invention relates to gene expression profiles, algorithms to generate gene expression profiles, microarrays comprising nucleic acid sequences representing gene expression profiles, methods of using gene expression profiles and microarrays, and business methods directed to the use of gene expression profiles, microarrays, and algorithms. The present invention further relates to protein expression profiles, algorithms to generate protein expression profiles, microarrays comprising protein-capture agents that bind proteins comprising protein expression profiles, methods of using protein expression profiles and microarrays, and business methods directed to the use of protein expression profiles, microarrays, and algorithms.

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US02/08456

A. CLASSIFICATION OF SUBJECT MATTER IPC(7) : C12Q 1/68 US CL : 435/6			
According to International Patent Classification (IPC) or to both national classification and IPC B. FIELDS SEARCHED			
Minimum documentation searched (classification system followed by classification symbols) U.S.: 435/6			
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched			
Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)			
C. DOCUMENTS CONSIDERED TO BE RELEVANT			
Category *	Citation of document, with indication, where a	opropriate, of the relevant passages	Relevant to claim No.
A, E	POOLE et al. Altered Patterns of Cellular Gene Exp Endothelial Cells Infected with Kaposi's Sarcoma-as Virology. April 2002, Vol 76, No. 7, pages 3395-34	sociated Herpesvius. Journal of	1, 3-4, 6-90
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A, E	US 6,403,316 B1 (SKALITER et al) 11 June 2002(11.06.2002), entire document.		1-90
Further	documents are listed in the continuation of Box C.	See patent family annex.	
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"P" document published prior to the international filing date but later than the priority date claimed		"&" document member of the same patent family	
Date of the actual completion of the international search		Date of mailing of the international search report	
30 September 2002 (30.09.2002)		06 NOV 2002	
	ailing address of the ISA/US missioner of Patents and Trademarks	Authorized officer	1 00
Box PCT		Shubo "Joe" Zhou Wella (allenash
Washington, D.C. 20231 Facsimile No. (703)305-3230		Telephone No. (703)-308-0196	